

SEQUENCE LISTING

<110> Pramod K. Srivastava

<120> ALPHA(2) MACROGLOBULIN RECEPTOR AS A HEAT SHOCK
PROTEIN RECEPTOR AND USES THEREOF

<130> 8449-123

<140> 09/625,137

<141> 2000-07-25

<150> 60/209,095

<151> 2000-06-02

<160> 59

<170> FastSEQ for Windows Version 3.0

<210> 1

<211> 14849

<212> DNA

<213> Mus musculus

<400> 1

cgctgctccc	cgccagtgca	ctgaggaggg	ggaaacgggg	gagcccctag	tgctccatca	60
ggccccctacc	aaggcacccc	catcggggtcc	acgcccccca	ccccccaccc	cgctcctcc	120
caattgtgca	tttttgcagc	cggagtcggc	tcogagatgg	ggctgtgagc	ttcgccctgg	180
gaggggggaga	ggagcgagga	gtaaagcagg	ggtgaagggt	tcgaatttgg	gggcaggggg	240
cgcccccgcg	tcagcaggcc	cttcccaggg	ggctcggaac	tgtaccattt	cacctatgcc	300
cctggttcgc	tttgcttaag	gaaggataag	atagaagagt	cggggagagg	aagataaagg	360
gggaccccc	aattgggggg	ggcgaggaca	agaagtaaca	ggaccagagg	gtgggggctg	420
ctgtttgcat	cggcccacac	catgctgacc	cgcgcgttgc	tgctgctcgt	gccgctgctt	480
tcagctctgg	tctccggggc	cactatggat	gccccataaa	cttgacagccc	taagcagttt	540
gcctgcagag	accaaatacac	ctgtatctca	aagggtctgg	ggtgtgacgg	tgaaagagat	600
tgccccgacg	gctctgatga	agccccctgag	atctgtccac	agagtaaagc	ccagagatgc	660
ccgccaatg	agcacagttg	tctggggact	gagctatgtg	tccccatgtc	tcgtctctgc	720
aacgggatcc	aggactgcat	ggatggctca	gacgaggggtg	ctcactgccg	agagctccga	780
gccaactgtt	ctcgaatggg	ttgtcaaac	cattgtgtac	ctacacccag	tgggcccacg	840
tgctactgta	acagcagctt	ccagctcgag	gcagatggca	agacgtgcaa	agattttgac	900
gagtgttccg	tgtatggcac	ctgcagccag	ctttgcacca	acacagatgg	ctccttcaca	960
tgtggctgtg	ttgaaggcta	cctgctgcaa	cgggacaacc	gctcctgcaa	ggccaagaat	1020
gagccagtag	atcgggccgcc	agtgtactg	attgccaaact	ctcagaacat	cctagctacg	1080
tacctgagtg	gggcccgaagt	gtctaccatc	acacccacca	gcacccgaca	aaccacggcc	1140
atggacttca	gttatgccaa	tgagaccgta	tgctgggtgc	acgttgggga	cagtgtctgcc	1200
cagacacagc	tcaagtgtgc	ccggatgcct	ggcctgaagg	gctttgtgga	tgagcatacc	1260
atcaacatct	ccctcagcct	gcaccacgtg	gagcagatgg	caatcgactg	gctgacggga	1320
aacttctact	ttgtcgacga	cattgacgac	aggatctttg	tctgtaaccg	aaacggggac	1380
acctgtgtca	ctctgctgga	cctggaactc	tacaacccca	aaggcatcgc	cttggaaccc	1440
gccatgggga	aggtgttctt	cactgactac	gggcagatcc	caaaggtgga	gcgctgtgac	1500
atggatggac	agaaccgcac	caagctgggtg	gatagcaaga	tcgtgtttcc	acacggcatc	1560
accctggacc	tggtcagccg	cctcgtctac	tgggcggacg	cctacctaga	ctacatcgag	1620
gtggtagact	acgaaggga	gggtcggcag	accatcatcc	aaggcatcct	gatcgagcac	1680
ctgtacggcc	tgaccgtgtt	tgagaactat	ctctacgcca	ccaactcgga	caatgccaac	1740
acgcagcaga	agacgagcgt	gatccgagtg	aaccgggttca	acagtactga	gtaccaggtc	1800
gtcaccctgtg	tggacaaggg	tggtgccctg	catatctacc	accagcgacg	ccagccccga	1860
gtgcggagtc	acgcctgtga	gaatgaccag	tacgggaagc	caggtggctg	ctccgacatc	1920
tgctcctctg	ccaacagtca	caaggcaagg	acctgcaggt	gcaggtctgg	cttcagcctg	1980
ggaagtgatg	ggaagtcttg	taagaaacct	gaacatgagc	tgttcctcgt	gtatggcaag	2040
ggccgaccag	gcatcattag	aggcatggac	atgggggcca	aggtcccaga	tgagcacatg	2100



atccccatcg	agaaccttat	gaatccacgc	gctctggact	tccacgccga	gaccggcttc	2160
atctactttg	ctgacaccac	cagctacctc	attggccgcc	agaaaattga	tggcacggag	2220
agagagacta	tcttgaagga	tggcatccac	aattgtggagg	gcgtagccgt	ggactggatg	2280
ggagacaatc	tttactggac	tgatgatggc	cccaagaaga	ccattagtgt	ggccaggctg	2340
gagaaagccg	ctcagacccg	gaagactcta	attgagggca	agatgacaca	ccccagggcc	2400
attgtagtgg	atccactcaa	tgggtggatg	tactggacag	actgggagga	ggaccccaag	2460
gacagtccgc	gagggcggct	cgagagggct	tggatggacg	gctcacaccg	agatatcttt	2520
gtcacctcca	agacagtgct	ttggcccaat	gggctaagcc	tggatatccc	agccggacgc	2580
ctctactggg	tggatgcctt	ctatgaccga	attgagacca	tactgctcaa	tggcacagac	2640
cggaaagattg	tatatgaggg	tcctgaactg	aatcatgcct	tcggcctgtg	tcaccatggc	2700
aactacctct	tttggaccga	gtaccggagc	ggcagcgtct	accgcttga	acggggcgtg	2760
gcaggcgcac	cgcccactgt	gaccttctg	cgcagcgaga	gaccgcctat	ctttgagatc	2820
cgaatgtacg	acgcgcacga	gcagcaagtg	ggtaaccaac	aatgccgggt	aaataacgga	2880
ggctgcagca	gcctgtgcct	cgccaccccc	gggagccgcc	agtgtgcctg	tgccgaggac	2940
caggtgttgg	acacagatgg	tgtcacctgc	ttggcgaacc	catcctacgt	gccccacccc	3000
cagtgccagc	cgggccagtt	tgcctgtgcc	aacaaccgct	gcattccagga	gcgctggaag	3060
tgtgacggag	acaacgactg	tctggacaac	agcgatgagg	ccccagcact	gtgccatcaa	3120
cacacctgtc	cctcggaccg	attcaagtgt	gagaacaacc	ggtgtatccc	caaccgctgg	3180
ctctgtgatg	gggataatga	ttgtggcaac	agcagggagc	aatccaatgc	cacgtgtctc	3240
gcccgcacct	gtccacccaa	ccagttctcc	tgtgccagtg	gccgatgcat	tcctatctca	3300
tggacctgtg	atctggatga	tgactgtggg	gaccggtccg	atgagtcaag	ctcatgcgcc	3360
tacccccact	gcttccccct	gactcaattt	acctgcaaca	atggcagatg	tattaacatc	3420
aactggcggg	gtgacaacga	caatgactgt	ggggacaaca	gcgacgaagc	cggctgcagt	3480
cactcctgct	ccagtaccca	gttcaagtgc	aacagtggca	gatgcatccc	cgagcactgg	3540
acgtgtgatg	gggacaatga	ttgtggggac	tacagcgacg	agacacacgc	caactgtacc	3600
aaccaggcta	caagacctcc	tgggtggctgc	cactcggatg	agttccagtg	cccgcctagat	3660
ggcctgtgca	tccccctgag	gtggcgctgc	gacggggaca	ccgactgcat	ggattccagc	3720
gatgagaaga	gctgtgaggg	cgtgacccat	gtttgtgacc	cgaatgtcaa	gtttggctgc	3780
aaggactccg	cccgtgcat	cagcaaggcg	tgggtgtgtg	atggcgacag	cgactgtgaa	3840
gataactccg	acgaggagaa	ctgtgaggcc	ctggcctgca	ggccaacctc	ccatccctgc	3900
gccaacaaca	cctctgtctg	cctgcctcct	gacaagctgt	gcgacggcaa	ggatgactgt	3960
ggagacggct	cggatgaggg	cgagctctgt	gaccagtgtt	ctctgaataa	tgggtggctgt	4020
agtcacaact	gctcagtggc	ccctggtgaa	ggcatcgtgt	gctcttgccc	tctgggcatg	4080
gagctgggct	ctgacaacca	cacctgccag	atccagagct	actgtgccaa	gcacctcaaa	4140
tgcagccaga	agtgtgacca	gaacaagttc	agtgtgaagt	gctcctgcta	cgagggctgg	4200
gtcttggagc	ctgacgggga	aacgtgccgc	agtctggatc	ccttcaaact	gttcatcatc	4260
ttctccaacc	gccacgagat	caggcgcatt	gaccttcaca	agggggacta	cagcgtccta	4320
gtgcctggcc	tgcgcaacac	tattgccctg	gacttccacc	tcagccagag	tgccctctac	4380
tggaccgcag	cggtagagga	caagatctac	cgtgggaaac	tcttggacaa	cggagccctg	4440
accagctttg	aggtggtgat	tcagtatggc	ttggccacac	cagagggcct	ggctgtagat	4500
tggattgcag	gcaacatcta	ctgggtggag	agcaacctgg	accagatcga	agtggccaag	4560
ctggacggaa	ccctccgaac	cactctgctg	gcgggtgaca	ttgagcacc	gagggccatc	4620
gctctggacc	ctcgggatgg	gattctgttt	tggacagact	gggatgccag	cctgccacga	4680
atcgaggctg	catccatgag	tggagctggc	cgccgaacca	tccaccggga	gacaggctct	4740
gggggctgcg	ccaatgggct	caccgtggat	tacctggaga	agcgcctcct	ctggattgat	4800
gctaggtcag	atgccatcta	ttcagcccg	tatgacggct	ccggccacat	ggaggtgctt	4860
cggggacacg	agttcctgtc	acacccattt	gccgtgacac	tgtacggtgg	ggaggtgtac	4920
tggaccgact	ggcgaacaaa	tacactggct	aaggccaaca	agtggactgg	ccacaacgtc	4980
accgtggtac	agaggaccaa	caccagccc	ttcgacctgc	aggtgtatca	cccttcccgg	5040
cagcccatgg	ctccaaaccc	atgtgaggcc	aatggcggcc	ggggcccctg	ttcccatctg	5100
tgcctcatca	actacaaccg	gaccgtctcc	tgggcctgtc	cccacctcat	gaagctgcac	5160
aaggacaaca	ccacctgcta	tgagtttaag	aagttcctgc	tgtacgcacg	tcagatggag	5220
atccggggcg	tggacctgga	tgccccgtac	tacaattata	tcatctcctt	cacgggtgct	5280
gatatcgaca	atgtcacggg	gctggactat	gatgcccag	agcagcgagt	ttactggctt	5340
gatgtgcgga	ctcaaggcca	caaaaaggca	tttatcaacg	gcactggcgt	ggagaccgtt	5400
gtctctgcag	acttgcccaa	cgcccacggg	ctggctgtgg	actgggtctc	ccgaaatctg	5460
ttttggacaa	gttacgacac	caacaagaag	cagattaacg	tggcccggct	ggacggctcc	5520
ttcaagaatg	cgggtggtgca	gggcctggag	cagccccacg	gcctggctgt	ccacccgctt	5580
cgtggcaagc	tctactggac	tgatggggac	aacatcagca	tggccaacat	ggatggggagc	5640
aaccacactc	tgtctctcag	tggccagaag	ggcctgtgtg	ggttggccat	tgacttcctt	5700
gagagcaaac	tctactggat	cagctctggg	aaccacacaa	tcaaccgttg	caatctggat	5760

gggagcgagc	tggaggtcat	cgacaccatg	cggagccagc	tgggcaaggc	cactgccctg	5820
gccatcatgg	gggacaagct	gtggtgggca	gatcaggtgt	cagagaagat	gggcacgtgc	5880
aacaaagccg	atggctctgg	gtccgtgggtg	ctgcggaaca	gtaccacgtt	ggttatgcac	5940
atgaaggtgt	atgacgagag	catccagcta	gagcatgagg	gcaccaaccc	ctgcagtgtc	6000
aacaacggag	actgttccca	gctctgectg	ccaacatcag	agacgactcg	ctcctgtatg	6060
tgtacagccg	gttacagcct	ccggagcgga	cagcaggcct	gtgaggggtg	gggctctttt	6120
ctcctgtact	ctgtacatga	gggaattcgg	gggattccac	tagatcccaa	tgacaagtgc	6180
gatgccctgg	tcccagtgtc	cggaacttca	ctggctgtcg	gaatcgactt	ccatgccgaa	6240
aatgacacta	tttattgggt	ggatatgggc	ctaagcacca	tcagcagggc	caagcgtgac	6300
cagacatggc	gagaggatgt	ggtgaccaac	ggtattggcc	gtgtggaggg	catcgccgtg	6360
gactggatcg	caggcaacat	atactggacg	gaccagggtc	tcgatgtcat	cgaggtggcc	6420
cggtccaatg	gctcttttcg	ttatgtggtc	atttcccagg	gtctggacaa	gccttgggccc	6480
atcactgtcc	accagagaaa	gggtacttgc	ttctggaccg	agtggggtca	ttaccacagt	6540
attgagcggg	ctcgccctga	tggcacagag	agagtgggtg	tggttaatgt	cagcatcagc	6600
tggcccaatg	gcatctcagt	agactatcag	ggcggcaagc	tctactgggtg	tgatgctcgg	6660
atggacaaga	tcgagcgcct	cgacctggaa	acgggcgaga	accgggaggt	ggtcctgtcc	6720
agcaataaca	tggatatgtt	ctccgtgtcc	gtgtttgagg	acttcatcta	ctggagtgc	6780
agaactcacg	ccaatggctc	catcaagcgc	ggctgcaaag	acaatgctac	agactccgtg	6840
cctctgagga	caggcattgg	tgttcagctt	aaagacatca	aggtcttcaa	cagggacagg	6900
cagaagggtg	ccaatgtgtg	cgcggtagcc	aacggcggtg	gccagcagct	ctgcttgtat	6960
cggggtggcg	gacagcgagc	ctgtgcctgt	gcccacggga	tgctggcaga	agacggggccc	7020
tcattgccgag	agtacgctgg	ctacctgtct	tactcagagc	ggaccatcct	caagagcatc	7080
cacctgtcgg	atgagcgtaa	cctcaacgca	ccggtgcagc	cctttgaaga	ccccgagcac	7140
atgaaaaatg	tcattgcctc	ggcctttgac	taccgagcag	gcacctcccc	ggggacccct	7200
aaccgcatct	tcttcagtga	catccacttt	gggaacatcc	agcagatcaa	tgacgatggc	7260
tcgggcagga	ccaccatcgt	ggaaaatgtg	ggctctgtgg	aaggcctggc	ctatcacctg	7320
ggctgggaca	cactgtactg	gacaagctac	accacatcca	ccatcacccg	ccacaccgtg	7380
gaccagactc	gcccaggggc	cttcgagagg	gagacagtca	tcaccatgtc	cggagacgac	7440
caccgcagag	cctttgtgct	ggatgagtgc	cagaacctga	tgttctggac	caattggaac	7500
gagctccatc	caagcatcat	gcgggcagcc	ctatccggag	ccaacgtcct	gacctcatt	7560
gagaaggaca	tcgcacgccc	caatgggttg	gccatcgacc	accgggcgga	gaagctgtac	7620
ttctcggatg	ccaccttgga	caagatcgag	cgctgcgagt	acgacggctc	ccaccgctat	7680
gtgatcctaa	agtcggagcc	cgtccacccc	tttgggttgg	cggtgtacgg	agagcacatt	7740
ttctggactg	actgggtgcg	gcgggctgtg	cagcgagcca	acaagtatgt	gggcagcgac	7800
atgaagctgc	ttcgggtgga	cattccccag	caacccatgg	gcatcatcgc	cgtggccaat	7860
gacaccaaca	gctgtgaact	ctccccctgc	cgtatcaaca	atggaggctg	ccaggatctg	7920
tgtctgctca	cccaccaagg	ccacgtcaac	tgttctgtgc	gagggggccg	gatcctcag	7980
gaggacttca	cctgcggggc	tgtgaactcc	tcttgcggg	cacaagatga	gtttgagtgt	8040
gccaatgggg	aatgtatcag	cttcagcctc	acctgtgatg	gcgtctccca	ctgcaaggac	8100
aagtccgatg	agaagccctc	ctactgcaac	tcacgccgct	gcaagaagac	tttccgccag	8160
tgtacaatg	gccgctgtgt	atccaacatg	ctgtggtgca	atggggtgga	ttactgtggg	8220
gatggctctg	atgagatacc	ttgcaacaag	actgcctgtg	gtgtgggtga	gttccgctgc	8280
cgggatgggt	cctgcacatg	gaactccagt	cgctgcaacc	agtttgtgga	ttgtgaggat	8340
gcctcggatg	agatgaattg	cagtgccaca	gactgcagca	gctatttccg	cctgggcgtg	8400
aaaggtgtcc	tcttcagacc	gtgcgagcgg	acatccctgt	gctacgcacc	tagctgggtg	8460
tgtgatggcg	ccaacgactg	tggagactac	agcgatgaac	gtgactgtcc	aggtgtgaag	8520
cgccctaggt	gcccgtcaa	ttactttgcc	tgccccagcg	ggcgctgtat	ccccatgagc	8580
tggacgtgtg	acaaggagga	tgactgtgag	aacggcgagg	atgagaccca	ctgcaacaag	8640
ttctgctcag	aggcacagtt	cgagtgccag	aaccaccggt	gtatctccaa	gcagtggctg	8700
tgtgacggta	gcgatgattg	cgggatggc	tccgatgagg	cagctcactg	tgaaggcaag	8760
acatgtggcc	cctcctcctt	ctcctgtccc	ggcaccacag	tgtgtgtccc	tgagcgctgg	8820
ctctgtgatg	gcgacaagga	ctgtaccgat	ggcgcggtatg	agagtgtcac	tgctggctgc	8880
ctgtacaaca	gcacctgtga	tgacctgtag	ttcatgtgcc	agaaccgctt	gtgtattccc	8940
aagcatttctg	tgtgcgacca	tgacctgtac	tgtgtgtgatg	gctctgatga	atccccctgag	9000
tgtgagtacc	caacctgcgg	gcccgaatgaa	ttccgctgtg	ccaatgggcg	ttgtctgagc	9060
tcccgtcagt	gggaatgtga	tggggagaa	gactgtcacg	accacagcga	tgaggctccc	9120
aagaaccac	actgcaccag	cccagagcac	aaatgcaatg	cctcatcaca	gttccctgtgc	9180
agcagcgggc	gctgcgtggc	tgaggcggtg	ctctgcaacg	gccaggacga	ctgtggggac	9240
ggttcagacg	aacgcgggtg	ccatgtcaac	gagtgtctca	gccgcaagct	cagtggctgc	9300
agtcaggact	gcgaggacct	caagataggc	tttaagtgcc	gctgtcgccc	gggcttccgg	9360
ctaaaggacg	atggcaggac	ctgtgccgac	ctggatgagt	gcagcaccac	cttccccctgc	9420

agccagctct	gcatacaac	ccacggaagt	tacaagtgtc	tgtgtgtgga	gggctatgca	9480
ccccgtggcg	gtgaccccc	cagctgcaaa	gctgtgaccg	atgaggagcc	atttctcatc	9540
tttgccaacc	gggtactacct	gcggaagctc	aacctggacg	gctccaacta	cacactgctt	9600
aagcagggcc	tgaacaatgc	ggtcgccttg	gcatttgact	accgagagca	gatgatctac	9660
tggacggggc	tgaccaccca	gggcagcatg	attcgcagga	tgcacctcaa	cggcagcaac	9720
gtgcagggtt	tgcaccggac	gggccttagt	aaccagatg	ggctcgctgt	ggactgggtg	9780
ggtggcaacc	tgtactggtg	tgacaagggc	agagatacca	ttgaggtgtc	caagcttaac	9840
ggggcctatc	ggacagtgtc	ggtcagctct	ggcctccggg	agcccagagc	tctggtagt	9900
gatgtacaga	atgggtacct	gtactggaca	gactgggggtg	accactcact	gatcgggccg	9960
attggcatgg	atggatctgg	ccgcagcatc	atcgtggaca	ctaagatcac	atggcccaat	10020
ggcctgaccg	tggactacgt	cacggaacgc	atctactggg	ctgacgcccg	tgaggactac	10080
atcgagtctg	ccagcctgga	tggctccaac	cgtcacgttg	tgctgagcca	agacatccca	10140
cacatctttg	cgctgacct	atttgaagac	tacgtctact	ggacagactg	ggaaacgaag	10200
tccatcaacc	gggcccacaa	gaccacgggt	gccaacaaaa	cactcctcat	cagcaccttg	10260
caccggccca	tggacttaca	tgtattccac	gccctgcgcc	agccagatgt	gcccacatcac	10320
ccctgcaaag	tcaacaatgg	tggctgcagc	aacctgtgcc	tgctgtcccc	tgggggtggt	10380
cacaagtgcg	cctgccccac	caacttctat	ctgggtggcg	atggccgtac	ctgtgtgtcc	10440
aactgcacag	caagccagtt	tgtgtgcaaa	aatgacaagt	gcatccccctt	ctggtggaag	10500
tgtgacacgg	aggacgactg	tggggatcac	tcagacgagc	ctccagactg	tcccaggttc	10560
aagtgcggcc	caggccagtt	ccagtgtctc	accggcatct	gcaccaaccc	tgccttcac	10620
tgtgatggcg	acaatgactg	ccaagacaat	agtgcagagg	ccaattgcga	cattcacgtc	10680
tgcttgccca	gccaatcaaa	gtgcaccaac	accaaccgct	gcattcctgg	catcttccgt	10740
tgcaatgggc	aggacaactg	cggggacggc	gaggatgagc	gggattgccc	tgaggtgacc	10800
tgcccccaca	accagttcca	gtgtccatc	accaagcgt	gcatccctcg	cgtctgggtc	10860
tgtgacaggg	ataatcactg	tgtggacggc	agtgatgagc	ctgccaaactg	tacccaaatg	10920
acctgtggag	tggatgagtt	ccgctgcaag	gattctggcc	gctgcatccc	cgcgcgctgg	10980
aagtgtgacg	gagaagatga	ctgtggggat	ggttcagatg	agcccaagga	agagtgtgat	11040
gagcgcacct	gtgagccata	ccagttccgc	tgcaaaaaa	accgctgtgt	cccaggccgt	11100
tggcaatgtg	actacgacaa	cgactgcgga	gataactcgg	acgaggagag	ctgcacacct	11160
cggccctgtg	ctgagagtga	gtttttctgt	gccaatggcc	gctgcatcgc	tgggcgctgg	11220
aagtgtgatg	gggaccatga	ctgtgccgac	ggctcagacg	agaaagactg	cacccccgc	11280
tgtgatattg	accagttcca	gtgcaagagt	ggccactgca	tccccctgcg	ctggccgtgt	11340
gacgcggatg	ctgactgtat	ggacggcagt	gacgaggaag	cctgtggcac	tggggtgagg	11400
acctgccccat	tggatgagtt	tcaatgtaac	aacaccttgt	gcaagccgct	ggcctggaag	11460
tgtgatggag	aggacgactg	tggggacaa	tcagatgaga	accccagagga	atgcgcccgg	11520
ttcatctgcc	ctcccaaccg	gcctttccgc	tgcaagaatg	accgagtctg	cctgtggatt	11580
gggcgccagt	gtgatggcgt	ggacaactgt	ggagatggga	ctgacgagga	ggactgtgag	11640
ccccccacgg	cccagaaccc	ccactgcaaa	gacaagaagg	agttcctgtg	ccgaaaccag	11700
cgtgtctctat	catcctccct	gcgtgttaac	atgttcgatg	actgcggcga	tggctccgat	11760
gaagaagatt	gcagcatcga	ccccaaagctg	accagctgtg	ccaccaatgc	cagcatgtgt	11820
ggggacgaag	ctcgttgtgt	gcgcactgag	aaagctgcct	actgtgcctg	ccgctcgggc	11880
ttccatactg	tgcggggcca	gcccggatgc	caggacatca	acgagtgcct	gcgctttggt	11940
acctgtctctc	agctctggaa	caaaccacaa	ggaggccacc	tctgcagctg	tgcccgcaac	12000
ttcatgaaga	cacacaacac	ctgcaaagct	gaaggctccg	agtaccaggt	gctatacatc	12060
gcgatgaca	acgagatccg	cagcttggtc	ccggggccacc	cccactcagc	ctacgagcag	12120
acattccagg	gcgatgagag	tgtccgcata	gatgccatgg	atgtccatgt	caaggccggc	12180
cgtgtctact	ggactaaactg	gcacacgggc	acaactctct	acaggagcct	gccccctgcc	12240
gccccctcta	ccacttccaa	ccgccaccgg	aggcagatcg	accgggggtgt	caccacctc	12300
aatatttcag	ggctgaagat	gccgaggggt	atcgtctatc	actgggtggc	cgggaatgtg	12360
tactggaccg	attccggccg	agacgtgatt	gaggtggcgc	aaatgaagg	cgagaaccgc	12420
aagacgctca	tctcgggcat	gattgatgag	ccccatgcca	tcgtgggtgga	ccctctgagg	12480
ggcaccatgt	actggtcaga	ctgggggaac	caccccaaga	ttgaaacagc	agcgatggat	12540
ggcacccttc	gggagactct	cgtgcaagac	aacattcagt	ggcctacagg	gctggctgtg	12600
gactatcaca	atgaacggct	ctactgggca	gatgccaaagc	tttcgggtcat	cggcagcatc	12660
cggctcaacg	gcactgacc	cattgtggct	gctgacagca	aacgaggcct	aagtcacccc	12720
ttcagcatcg	atgtgtttga	agactacatc	tacggagtca	cttacatcaa	taatcgtgtc	12780
ttcaagatcc	acaagtttgg	acacagcccc	ttgtacaacc	taactggggg	cctgagccat	12840
gcctctgatg	tagtccttta	ccatcaacac	aagcagcctg	aagtgaccaa	cccctgtgac	12900
cgcaagaaat	gcgaatggct	gtgtctgctg	agccccagcg	ggcctgtctg	cacctgtccc	12960
aatggaaaga	ggctggataa	tggcacctgt	gtgcctgtgc	cctctccaac	acccccctcca	13020
gatgccccta	ggcctggaac	ctgcactctg	cagtgtctca	atgggtggtag	ttgtttcctc	13080

aacgctcggg	ggcagcccaa	gtgccgttgc	cagccccgtt	acacaggcga	taagtgtgag	13140
ctggatcagt	gctgggaata	ctgtcacaac	ggaggcacct	gtgcggcttc	cccatctggc	13200
atgcccacgt	gccgctgtcc	cactggcttc	acgggcccc	aatgcaccgc	acagggtgtg	13260
gcaggctact	gctctaacaa	cagcacctgc	accgtcaacc	agggaacca	gccccagtgc	13320
cgatgtctac	ctggcttcct	gggcgaccgt	tgccagtacc	ggcagtgtct	tggcttctgt	13380
gagaactttg	gcacctgtca	gatggctgct	gatggctccc	gacaatgtcg	ctgcaccgtc	13440
tactttgagg	gaccaaggtg	tgaggtgaac	aagtgtagtc	gctgtctcca	aggcgcctgt	13500
gtggtcaata	agcagaccgg	agatgtcaca	tgcaactgca	ctgatggccg	ggtagcccc	13560
agttgtctca	cctgcacgca	tcactgtagc	aatggtggct	cctgcaccat	gaacagcaag	13620
atgatgcctg	agtgccagt	cccgcctcat	atgacaggac	cccgttgcca	ggagcaggtt	13680
gttagtcagc	aacagcctgg	gcataatggc	tccatcctga	tccctctgct	gctgcttctc	13740
ctgctgcttc	tggtggctgg	cgtggtgttc	tggtataagc	ggcgagtccg	aggggctaag	13800
ggcttccagc	accagcggtg	gaccaatggg	gccatgaatg	tggaaattgg	aaaccctacc	13860
tacaagatgt	atgaaggtgg	agagcccgat	gatgtcgggg	gcctactgga	tgctgatttt	13920
gccccttgacc	ctgacaagcc	taccaacttc	accaacccag	tgtatgccac	gctctacatg	13980
ggggggccacg	gcagccgcca	ttccctggcc	agcacggacg	agaagcgaga	actgctgggc	14040
cggggacctg	aagacgagat	aggagatccc	ttggcatagg	gccctgcccc	gacggatgtc	14100
cccagaaagc	cccctgccac	atgagtcttt	caatgaaccc	cctccccagc	cgcccttctt	14160
ccggccctgc	cggtgttaca	aatgtaaaaa	tgaaggaatt	actttttata	tgtgagcgag	14220
caagcgagca	agcacagtat	tatctctttg	catttctctc	ctgcctgctc	ctcagtatcc	14280
cccccatgct	gccttgaggg	ggcggggagg	gctttgtggc	tcaaagggtat	gaaggagtcc	14340
acatgttccc	taccgagcat	acccctggaa	gcctggcggc	acggcctccc	caccacgcct	14400
gtgcaagaca	ctcaacgggg	ctccgtgtcc	cagctttcct	ttccttggct	ctctgggggt	14460
agttcagggg	aggtggagtc	ctctgctgac	cctgtctgga	agatttggct	ctagctgagg	14520
aaggagtctt	ttagttgagg	gaagtcaccc	caaaccccag	ctcccacttt	caggggcacc	14580
tctcagatgg	ccatgctcag	tatcccttcc	agacaggccc	tccctctctt	agcgcctcct	14640
ctgtggctcc	tagggctgaa	cacattcttt	ggtaactgtc	ccccaaagcct	cccatccccc	14700
tgagggccag	gaagagtcgg	ggcacaccaa	ggaagggcaa	gcgggcagcc	ccattttggg	14760
gacgtgaacg	ttttaataat	ttttgctgaa	ttcctttaca	actaaataac	acagatattg	14820
ttataaataa	aattgtaaaa	aaaaaaaaa				14849

<210> 2
 <211> 4545
 <212> PRT
 <213> Mus musculus

<400> 2

Met	Leu	Thr	Pro	Pro	Leu	Leu	Leu	Leu	Val	Pro	Leu	Leu	Ser	Ala	Leu
1				5					10					15	
Val	Ser	Gly	Ala	Thr	Met	Asp	Ala	Pro	Lys	Thr	Cys	Ser	Pro	Lys	Gln
			20					25					30		
Phe	Ala	Cys	Arg	Asp	Gln	Ile	Thr	Cys	Ile	Ser	Lys	Gly	Trp	Arg	Cys
		35					40					45			
Asp	Gly	Glu	Arg	Asp	Cys	Pro	Asp	Gly	Ser	Asp	Glu	Ala	Pro	Glu	Ile
	50				55						60				
Cys	Pro	Gln	Ser	Lys	Ala	Gln	Arg	Cys	Pro	Pro	Asn	Glu	His	Ser	Cys
65					70					75				80	
Leu	Gly	Thr	Glu	Leu	Cys	Val	Pro	Met	Ser	Arg	Leu	Cys	Asn	Gly	Ile
			85					90					95		
Gln	Asp	Cys	Met	Asp	Gly	Ser	Asp	Glu	Gly	Ala	His	Cys	Arg	Glu	Leu
			100					105					110		
Arg	Ala	Asn	Cys	Ser	Arg	Met	Gly	Cys	Gln	His	His	Cys	Val	Pro	Thr
		115					120					125			
Pro	Ser	Gly	Pro	Thr	Cys	Tyr	Cys	Asn	Ser	Ser	Phe	Gln	Leu	Glu	Ala
		130				135					140				
Asp	Gly	Lys	Thr	Cys	Lys	Asp	Phe	Asp	Glu	Cys	Ser	Val	Tyr	Gly	Thr
145					150					155				160	
Cys	Ser	Gln	Leu	Cys	Thr	Asn	Thr	Asp	Gly	Ser	Phe	Thr	Cys	Gly	Cys
			165					170						175	

Val	Glu	Gly	Tyr	Leu	Leu	Gln	Pro	Asp	Asn	Arg	Ser	Cys	Lys	Ala	Lys
			180					185					190		
Asn	Glu	Pro	Val	Asp	Arg	Pro	Pro	Val	Leu	Leu	Ile	Ala	Asn	Ser	Gln
		195					200					205			
Asn	Ile	Leu	Ala	Thr	Tyr	Leu	Ser	Gly	Ala	Gln	Val	Ser	Thr	Ile	Thr
	210					215					220				
Pro	Thr	Ser	Thr	Arg	Gln	Thr	Thr	Ala	Met	Asp	Phe	Ser	Tyr	Ala	Asn
225					230					235					240
Glu	Thr	Val	Cys	Trp	Val	His	Val	Gly	Asp	Ser	Ala	Ala	Gln	Thr	Gln
			245						250					255	
Leu	Lys	Cys	Ala	Arg	Met	Pro	Gly	Leu	Lys	Gly	Phe	Val	Asp	Glu	His
		260						265					270		
Thr	Ile	Asn	Ile	Ser	Leu	Ser	Leu	His	His	Val	Glu	Gln	Met	Ala	Ile
		275					280						285		
Asp	Trp	Leu	Thr	Gly	Asn	Phe	Tyr	Phe	Val	Asp	Asp	Ile	Asp	Asp	Arg
	290				295						300				
Ile	Phe	Val	Cys	Asn	Arg	Asn	Gly	Asp	Thr	Cys	Val	Thr	Leu	Leu	Asp
305					310					315					320
Leu	Glu	Leu	Tyr	Asn	Pro	Lys	Gly	Ile	Ala	Leu	Asp	Pro	Ala	Met	Gly
			325						330					335	
Lys	Val	Phe	Phe	Thr	Asp	Tyr	Gly	Gln	Ile	Pro	Lys	Val	Glu	Arg	Cys
		340						345					350		
Asp	Met	Asp	Gly	Gln	Asn	Arg	Thr	Lys	Leu	Val	Asp	Ser	Lys	Ile	Val
		355					360					365			
Phe	Pro	His	Gly	Ile	Thr	Leu	Asp	Leu	Val	Ser	Arg	Leu	Val	Tyr	Trp
	370				375						380				
Ala	Asp	Ala	Tyr	Leu	Asp	Tyr	Ile	Glu	Val	Val	Asp	Tyr	Glu	Gly	Lys
385					390					395					400
Gly	Arg	Gln	Thr	Ile	Ile	Gln	Gly	Ile	Leu	Ile	Glu	His	Leu	Tyr	Gly
				405					410					415	
Leu	Thr	Val	Phe	Glu	Asn	Tyr	Leu	Tyr	Ala	Thr	Asn	Ser	Asp	Asn	Ala
		420						425					430		
Asn	Thr	Gln	Gln	Lys	Thr	Ser	Val	Ile	Arg	Val	Asn	Arg	Phe	Asn	Ser
		435					440					445			
Thr	Glu	Tyr	Gln	Val	Val	Thr	Arg	Val	Asp	Lys	Gly	Gly	Ala	Leu	His
	450					455					460				
Ile	Tyr	His	Gln	Arg	Arg	Gln	Pro	Arg	Val	Arg	Ser	His	Ala	Cys	Glu
465					470					475					480
Asn	Asp	Gln	Tyr	Gly	Lys	Pro	Gly	Gly	Cys	Ser	Asp	Ile	Cys	Leu	Leu
				485					490					495	
Ala	Asn	Ser	His	Lys	Ala	Arg	Thr	Cys	Arg	Cys	Arg	Ser	Gly	Phe	Ser
		500						505					510		
Leu	Gly	Ser	Asp	Gly	Lys	Ser	Cys	Lys	Lys	Pro	Glu	His	Glu	Leu	Phe
		515					520					525			
Leu	Val	Tyr	Gly	Lys	Gly	Arg	Pro	Gly	Ile	Ile	Arg	Gly	Met	Asp	Met
	530					535					540				
Gly	Ala	Lys	Val	Pro	Asp	Glu	His	Met	Ile	Pro	Ile	Glu	Asn	Leu	Met
545					550					555					560
Asn	Pro	Arg	Ala	Leu	Asp	Phe	His	Ala	Glu	Thr	Gly	Phe	Ile	Tyr	Phe
			565						570					575	
Ala	Asp	Thr	Thr	Ser	Tyr	Leu	Ile	Gly	Arg	Gln	Lys	Ile	Asp	Gly	Thr
		580						585					590		
Glu	Arg	Glu	Thr	Ile	Leu	Lys	Asp	Gly	Ile	His	Asn	Val	Glu	Gly	Val
	595						600					605			
Ala	Val	Asp	Trp	Met	Gly	Asp	Asn	Leu	Tyr	Trp	Thr	Asp	Asp	Gly	Pro
	610					615					620				
Lys	Lys	Thr	Ile	Ser	Val	Ala	Arg	Leu	Glu	Lys	Ala	Ala	Gln	Thr	Arg
625					630					635					640
Lys	Thr	Leu	Ile	Glu	Gly	Lys	Met	Thr	His	Pro	Arg	Ala	Ile	Val	Val
			645						650					655	
Asp	Pro	Leu	Asn	Gly	Trp	Met	Tyr	Trp	Thr	Asp	Trp	Glu	Glu	Asp	Pro

Cys Ala Asn Asn Thr Ser Val Cys Leu Pro Pro Asp Lys Leu Cys Asp
1155 1160 1165
Gly Lys Asp Asp Cys Gly Asp Gly Ser Asp Glu Gly Glu Leu Cys Asp
1170 1175 1180
Gln Cys Ser Leu Asn Asn Gly Gly Cys Ser His Asn Cys Ser Val Ala
1185 1190 1195 1200
Pro Gly Glu Gly Ile Val Cys Ser Cys Pro Leu Gly Met Glu Leu Gly
1205 1210 1215
Ser Asp Asn His Thr Cys Gln Ile Gln Ser Tyr Cys Ala Lys His Leu
1220 1225 1230
Lys Cys Ser Gln Lys Cys Asp Gln Asn Lys Phe Ser Val Lys Cys Ser
1235 1240 1245
Cys Tyr Glu Gly Trp Val Leu Glu Pro Asp Gly Glu Thr Cys Arg Ser
1250 1255 1260
Leu Asp Pro Phe Lys Leu Phe Ile Ile Phe Ser Asn Arg His Glu Ile
1265 1270 1275 1280
Arg Arg Ile Asp Leu His Lys Gly Asp Tyr Ser Val Leu Val Pro Gly
1285 1290 1295
Leu Arg Asn Thr Ile Ala Leu Asp Phe His Leu Ser Gln Ser Ala Leu
1300 1305 1310
Tyr Trp Thr Asp Ala Val Glu Asp Lys Ile Tyr Arg Gly Lys Leu Leu
1315 1320 1325
Asp Asn Gly Ala Leu Thr Ser Phe Glu Val Val Ile Gln Tyr Gly Leu
1330 1335 1340
Ala Thr Pro Glu Gly Leu Ala Val Asp Trp Ile Ala Gly Asn Ile Tyr
1345 1350 1355 1360
Trp Val Glu Ser Asn Leu Asp Gln Ile Glu Val Ala Lys Leu Asp Gly
1365 1370 1375
Thr Leu Arg Thr Thr Leu Leu Ala Gly Asp Ile Glu His Pro Arg Ala
1380 1385 1390
Ile Ala Leu Asp Pro Arg Asp Gly Ile Leu Phe Trp Thr Asp Trp Asp
1395 1400 1405
Ala Ser Leu Pro Arg Ile Glu Ala Ala Ser Met Ser Gly Ala Gly Arg
1410 1415 1420
Arg Thr Ile His Arg Glu Thr Gly Ser Gly Gly Cys Ala Asn Gly Leu
1425 1430 1435 1440
Thr Val Asp Tyr Leu Glu Lys Arg Ile Leu Trp Ile Asp Ala Arg Ser
1445 1450 1455
Asp Ala Ile Tyr Ser Ala Arg Tyr Asp Gly Ser Gly His Met Glu Val
1460 1465 1470
Leu Arg Gly His Glu Phe Leu Ser His Pro Phe Ala Val Thr Leu Tyr
1475 1480 1485
Gly Gly Glu Val Tyr Trp Thr Asp Trp Arg Thr Asn Thr Leu Ala Lys
1490 1495 1500
Ala Asn Lys Trp Thr Gly His Asn Val Thr Val Val Gln Arg Thr Asn
1505 1510 1515 1520
Thr Gln Pro Phe Asp Leu Gln Val Tyr His Pro Ser Arg Gln Pro Met
1525 1530 1535
Ala Pro Asn Pro Cys Glu Ala Asn Gly Gly Arg Gly Pro Cys Ser His
1540 1545 1550
Leu Cys Leu Ile Asn Tyr Asn Arg Thr Val Ser Trp Ala Cys Pro His
1555 1560 1565
Leu Met Lys Leu His Lys Asp Asn Thr Thr Cys Tyr Glu Phe Lys Lys
1570 1575 1580
Phe Leu Leu Tyr Ala Arg Gln Met Glu Ile Arg Gly Val Asp Leu Asp
1585 1590 1595 1600
Ala Pro Tyr Tyr Asn Tyr Ile Ile Ser Phe Thr Val Pro Asp Ile Asp
1605 1610 1615
Asn Val Thr Val Leu Asp Tyr Asp Ala Arg Glu Gln Arg Val Tyr Trp
1620 1625 1630
Ser Asp Val Arg Thr Gln Ala Ile Lys Arg Ala Phe Ile Asn Gly Thr

Ala Thr Asp Ser Val Pro Leu Arg Thr Gly Ile Gly Val Gln Leu Lys
 2130 2135 2140
 Asp Ile Lys Val Phe Asn Arg Asp Arg Gln Lys Gly Thr Asn Val Cys
 2145 2150 2155 2160
 Ala Val Ala Asn Gly Gly Cys Gln Gln Leu Cys Leu Tyr Arg Gly Gly
 2165 2170 2175
 Gly Gln Arg Ala Cys Ala Cys Ala His Gly Met Leu Ala Glu Asp Gly
 2180 2185 2190
 Ala Ser Cys Arg Glu Tyr Ala Gly Tyr Leu Leu Tyr Ser Glu Arg Thr
 2195 2200 2205
 Ile Leu Lys Ser Ile His Leu Ser Asp Glu Arg Asn Leu Asn Ala Pro
 2210 2215 2220
 Val Gln Pro Phe Glu Asp Pro Glu His Met Lys Asn Val Ile Ala Leu
 2225 2230 2235 2240
 Ala Phe Asp Tyr Arg Ala Gly Thr Ser Pro Gly Thr Pro Asn Arg Ile
 2245 2250 2255
 Phe Phe Ser Asp Ile His Phe Gly Asn Ile Gln Gln Ile Asn Asp Asp
 2260 2265 2270
 Gly Ser Gly Arg Thr Thr Ile Val Glu Asn Val Gly Ser Val Glu Gly
 2275 2280 2285
 Leu Ala Tyr His Arg Gly Trp Asp Thr Leu Tyr Trp Thr Ser Tyr Thr
 2290 2295 2300
 Thr Ser Thr Ile Thr Arg His Thr Val Asp Gln Thr Arg Pro Gly Ala
 2305 2310 2315 2320
 Phe Glu Arg Glu Thr Val Ile Thr Met Ser Gly Asp Asp His Pro Arg
 2325 2330 2335
 Ala Phe Val Leu Asp Glu Cys Gln Asn Leu Met Phe Trp Thr Asn Trp
 2340 2345 2350
 Asn Glu Leu His Pro Ser Ile Met Arg Ala Ala Leu Ser Gly Ala Asn
 2355 2360 2365
 Val Leu Thr Leu Ile Glu Lys Asp Ile Arg Thr Pro Asn Gly Leu Ala
 2370 2375 2380
 Ile Asp His Arg Ala Glu Lys Leu Tyr Phe Ser Asp Ala Thr Leu Asp
 2385 2390 2395 2400
 Lys Ile Glu Arg Cys Glu Tyr Asp Gly Ser His Arg Tyr Val Ile Leu
 2405 2410 2415
 Lys Ser Glu Pro Val His Pro Phe Gly Leu Ala Val Tyr Gly Glu His
 2420 2425 2430
 Ile Phe Trp Thr Asp Trp Val Arg Arg Ala Val Gln Arg Ala Asn Lys
 2435 2440 2445
 Tyr Val Gly Ser Asp Met Lys Leu Leu Arg Val Asp Ile Pro Gln Gln
 2450 2455 2460
 Pro Met Gly Ile Ile Ala Val Ala Asn Asp Thr Asn Ser Cys Glu Leu
 2465 2470 2475 2480
 Ser Pro Cys Arg Ile Asn Asn Gly Gly Cys Gln Asp Leu Cys Leu Leu
 2485 2490 2495
 Thr His Gln Gly His Val Asn Cys Ser Cys Arg Gly Gly Arg Ile Leu
 2500 2505 2510
 Gln Glu Asp Phe Thr Cys Arg Ala Val Asn Ser Ser Cys Arg Ala Gln
 2515 2520 2525
 Asp Glu Phe Glu Cys Ala Asn Gly Glu Cys Ile Ser Phe Ser Leu Thr
 2530 2535 2540
 Cys Asp Gly Val Ser His Cys Lys Asp Lys Ser Asp Glu Lys Pro Ser
 2545 2550 2555 2560
 Tyr Cys Asn Ser Arg Arg Cys Lys Lys Thr Phe Arg Gln Cys Asn Asn
 2565 2570 2575
 Gly Arg Cys Val Ser Asn Met Leu Trp Cys Asn Gly Val Asp Tyr Cys
 2580 2585 2590
 Gly Asp Gly Ser Asp Glu Ile Pro Cys Asn Lys Thr Ala Cys Gly Val
 2595 2600 2605
 Gly Glu Phe Arg Cys Arg Asp Gly Ser Cys Ile Gly Asn Ser Ser Arg

2610	2615	2620
Cys Asn Gln Phe Val Asp Cys Glu Asp Ala Ser Asp Glu Met Asn Cys		
2625	2630	2635
Ser Ala Thr Asp Cys Ser Ser Tyr Phe Arg Leu Gly Val Lys Gly Val		2640
	2645	2650
Leu Phe Gln Pro Cys Glu Arg Thr Ser Leu Cys Tyr Ala Pro Ser Trp		2655
	2660	2665
Val Cys Asp Gly Ala Asn Asp Cys Gly Asp Tyr Ser Asp Glu Arg Asp		2670
	2675	2680
Cys Pro Gly Val Lys Arg Pro Arg Cys Pro Leu Asn Tyr Phe Ala Cys		2685
	2690	2695
Pro Ser Gly Arg Cys Ile Pro Met Ser Trp Thr Cys Asp Lys Glu Asp		2700
2705	2710	2715
Asp Cys Glu Asn Gly Glu Asp Glu Thr His Cys Asn Lys Phe Cys Ser		2720
	2725	2730
Glu Ala Gln Phe Glu Cys Gln Asn His Arg Cys Ile Ser Lys Gln Trp		2735
	2740	2745
Leu Cys Asp Gly Ser Asp Asp Cys Gly Asp Gly Ser Asp Glu Ala Ala		2750
	2755	2760
His Cys Glu Gly Lys Thr Cys Gly Pro Ser Ser Phe Ser Cys Pro Gly		2765
	2770	2775
Thr His Val Cys Val Pro Glu Arg Trp Leu Cys Asp Gly Asp Lys Asp		2780
2785	2790	2795
Cys Thr Asp Gly Ala Asp Glu Ser Val Thr Ala Gly Cys Leu Tyr Asn		2800
	2805	2810
Ser Thr Cys Asp Asp Arg Glu Phe Met Cys Gln Asn Arg Leu Cys Ile		2815
	2820	2825
Pro Lys His Phe Val Cys Asp His Asp Arg Asp Cys Ala Asp Gly Ser		2830
	2835	2840
Asp Glu Ser Pro Glu Cys Glu Tyr Pro Thr Cys Gly Pro Asn Glu Phe		2845
	2850	2855
Arg Cys Ala Asn Gly Arg Cys Leu Ser Ser Arg Gln Trp Glu Cys Asp		2860
2865	2870	2875
Gly Glu Asn Asp Cys His Asp His Ser Asp Glu Ala Pro Lys Asn Pro		2880
	2885	2890
His Cys Thr Ser Pro Glu His Lys Cys Asn Ala Ser Ser Gln Phe Leu		2895
	2900	2905
Cys Ser Ser Gly Arg Cys Val Ala Glu Ala Leu Leu Cys Asn Gly Gln		2910
	2915	2920
Asp Asp Cys Gly Asp Gly Ser Asp Glu Arg Gly Cys His Val Asn Glu		2925
	2930	2935
Cys Leu Ser Arg Lys Leu Ser Gly Cys Ser Gln Asp Cys Glu Asp Leu		2940
2945	2950	2955
Lys Ile Gly Phe Lys Cys Arg Cys Arg Pro Gly Phe Arg Leu Lys Asp		2960
	2965	2970
Asp Gly Arg Thr Cys Ala Asp Leu Asp Glu Cys Ser Thr Thr Phe Pro		2975
	2980	2985
Cys Ser Gln Leu Cys Ile Asn Thr His Gly Ser Tyr Lys Cys Leu Cys		2990
	2995	3000
Val Glu Gly Tyr Ala Pro Arg Gly Gly Asp Pro His Ser Cys Lys Ala		3005
	3010	3015
Val Thr Asp Glu Glu Pro Phe Leu Ile Phe Ala Asn Arg Tyr Tyr Leu		3020
3025	3030	3035
Arg Lys Leu Asn Leu Asp Gly Ser Asn Tyr Thr Leu Leu Lys Gln Gly		3040
	3045	3050
Leu Asn Asn Ala Val Ala Leu Ala Phe Asp Tyr Arg Glu Gln Met Ile		3055
	3060	3065
Tyr Trp Thr Gly Val Thr Thr Gln Gly Ser Met Ile Arg Arg Met His		3070
	3075	3080
Leu Asn Gly Ser Asn Val Gln Val Leu His Arg Thr Gly Leu Ser Asn		3085
	3090	3095
		3100

Pro Asp Gly Leu Ala Val Asp Trp Val Gly Gly Asn Leu Tyr Trp Cys
 3105 3110 3115 3120
 Asp Lys Gly Arg Asp Thr Ile Glu Val Ser Lys Leu Asn Gly Ala Tyr
 3125 3130 3135
 Arg Thr Val Leu Val Ser Ser Gly Leu Arg Glu Pro Arg Ala Leu Val
 3140 3145 3150
 Val Asp Val Gln Asn Gly Tyr Leu Tyr Trp Thr Asp Trp Gly Asp His
 3155 3160 3165
 Ser Leu Ile Gly Arg Ile Gly Met Asp Gly Ser Gly Arg Ser Ile Ile
 3170 3175 3180
 Val Asp Thr Lys Ile Thr Trp Pro Asn Gly Leu Thr Val Asp Tyr Val
 3185 3190 3195 3200
 Thr Glu Arg Ile Tyr Trp Ala Asp Ala Arg Glu Asp Tyr Ile Glu Phe
 3205 3210 3215
 Ala Ser Leu Asp Gly Ser Asn Arg His Val Val Leu Ser Gln Asp Ile
 3220 3225 3230
 Pro His Ile Phe Ala Leu Thr Leu Phe Glu Asp Tyr Val Tyr Trp Thr
 3235 3240 3245
 Asp Trp Glu Thr Lys Ser Ile Asn Arg Ala His Lys Thr Thr Gly Ala
 3250 3255 3260
 Asn Lys Thr Leu Leu Ile Ser Thr Leu His Arg Pro Met Asp Leu His
 3265 3270 3275 3280
 Val Phe His Ala Leu Arg Gln Pro Asp Val Pro Asn His Pro Cys Lys
 3285 3290 3295
 Val Asn Asn Gly Gly Cys Ser Asn Leu Cys Leu Leu Ser Pro Gly Gly
 3300 3305 3310
 Gly His Lys Cys Ala Cys Pro Thr Asn Phe Tyr Leu Gly Gly Asp Gly
 3315 3320 3325
 Arg Thr Cys Val Ser Asn Cys Thr Ala Ser Gln Phe Val Cys Lys Asn
 3330 3335 3340
 Asp Lys Cys Ile Pro Phe Trp Trp Lys Cys Asp Thr Glu Asp Asp Cys
 3345 3350 3355 3360
 Gly Asp His Ser Asp Glu Pro Pro Asp Cys Pro Glu Phe Lys Cys Arg
 3365 3370 3375
 Pro Gly Gln Phe Gln Cys Ser Thr Gly Ile Cys Thr Asn Pro Ala Phe
 3380 3385 3390
 Ile Cys Asp Gly Asp Asn Asp Cys Gln Asp Asn Ser Asp Glu Ala Asn
 3395 3400 3405
 Cys Asp Ile His Val Cys Leu Pro Ser Gln Phe Lys Cys Thr Asn Thr
 3410 3415 3420
 Asn Arg Cys Ile Pro Gly Ile Phe Arg Cys Asn Gly Gln Asp Asn Cys
 3425 3430 3435 3440
 Gly Asp Gly Glu Asp Glu Arg Asp Cys Pro Glu Val Thr Cys Ala Pro
 3445 3450 3455
 Asn Gln Phe Gln Cys Ser Ile Thr Lys Arg Cys Ile Pro Arg Val Trp
 3460 3465 3470
 Val Cys Asp Arg Asp Asn His Cys Val Asp Gly Ser Asp Glu Pro Ala
 3475 3480 3485
 Asn Cys Thr Gln Met Thr Cys Gly Val Asp Glu Phe Arg Cys Lys Asp
 3490 3495 3500
 Ser Gly Arg Cys Ile Pro Ala Arg Trp Lys Cys Asp Gly Glu Asp Asp
 3505 3510 3515 3520
 Cys Gly Asp Gly Ser Asp Glu Pro Lys Glu Glu Cys Asp Glu Arg Thr
 3525 3530 3535
 Cys Glu Pro Tyr Gln Phe Arg Cys Lys Asn Asn Arg Cys Val Pro Gly
 3540 3545 3550
 Arg Trp Gln Cys Asp Tyr Asp Asn Asp Cys Gly Asp Asn Ser Asp Glu
 3555 3560 3565
 Glu Ser Cys Thr Pro Arg Pro Cys Ser Glu Ser Glu Phe Phe Cys Ala
 3570 3575 3580
 Asn Gly Arg Cys Ile Ala Gly Arg Trp Lys Cys Asp Gly Asp His Asp

3585		3590		3595		3600
Cys Ala Asp Gly Ser	Asp Glu Lys Asp Cys Thr Pro Arg Cys Asp Met					
	3605		3610			3615
Asp Gln Phe Gln Cys Lys Ser Gly His Cys Ile Pro Leu Arg Trp Pro						
	3620		3625			3630
Cys Asp Ala Asp Ala Asp Cys Met Asp Gly Ser Asp Glu Glu Ala Cys						
	3635		3640			3645
Gly Thr Gly Val Arg Thr Cys Pro Leu Asp Glu Phe Gln Cys Asn Asn						
	3650		3655			3660
Thr Leu Cys Lys Pro Leu Ala Trp Lys Cys Asp Gly Glu Asp Asp Cys						
	3665		3670			3675
Gly Asp Asn Ser Asp Glu Asn Pro Glu Glu Cys Ala Arg Phe Ile Cys						
	3685		3690			3695
Pro Pro Asn Arg Pro Phe Arg Cys Lys Asn Asp Arg Val Cys Leu Trp						
	3700		3705			3710
Ile Gly Arg Gln Cys Asp Gly Val Asp Asn Cys Gly Asp Gly Thr Asp						
	3715		3720			3725
Glu Glu Asp Cys Glu Pro Pro Thr Ala Gln Asn Pro His Cys Lys Asp						
	3730		3735			3740
Lys Lys Glu Phe Leu Cys Arg Asn Gln Arg Cys Leu Ser Ser Ser Leu						
	3745		3750			3755
Arg Cys Asn Met Phe Asp Asp Cys Gly Asp Gly Ser Asp Glu Glu Asp						
	3765		3770			3775
Cys Ser Ile Asp Pro Lys Leu Thr Ser Cys Ala Thr Asn Ala Ser Met						
	3780		3785			3790
Cys Gly Asp Glu Ala Arg Cys Val Arg Thr Glu Lys Ala Ala Tyr Cys						
	3795		3800			3805
Ala Cys Arg Ser Gly Phe His Thr Val Pro Gly Gln Pro Gly Cys Gln						
	3810		3815			3820
Asp Ile Asn Glu Cys Leu Arg Phe Gly Thr Cys Ser Gln Leu Trp Asn						
	3825		3830			3835
Lys Pro Lys Gly Gly His Leu Cys Ser Cys Ala Arg Asn Phe Met Lys						
	3845		3850			3855
Thr His Asn Thr Cys Lys Ala Glu Gly Ser Glu Tyr Gln Val Leu Tyr						
	3860		3865			3870
Ile Ala Asp Asp Asn Glu Ile Arg Ser Leu Phe Pro Gly His Pro His						
	3875		3880			3885
Ser Ala Tyr Glu Gln Thr Phe Gln Gly Asp Glu Ser Val Arg Ile Asp						
	3890		3895			3900
Ala Met Asp Val His Val Lys Ala Gly Arg Val Tyr Trp Thr Asn Trp						
	3905		3910			3915
His Thr Gly Thr Ile Ser Tyr Arg Ser Leu Pro Pro Ala Ala Pro Pro						
	3925		3930			3935
Thr Thr Ser Asn Arg His Arg Arg Gln Ile Asp Arg Gly Val Thr His						
	3940		3945			3950
Leu Asn Ile Ser Gly Leu Lys Met Pro Arg Gly Ile Ala Ile Asp Trp						
	3955		3960			3965
Val Ala Gly Asn Val Tyr Trp Thr Asp Ser Gly Arg Asp Val Ile Glu						
	3970		3975			3980
Val Ala Gln Met Lys Gly Glu Asn Arg Lys Thr Leu Ile Ser Gly Met						
	3985		3990			3995
Ile Asp Glu Pro His Ala Ile Val Val Asp Pro Leu Arg Gly Thr Met						
	4005		4010			4015
Tyr Trp Ser Asp Trp Gly Asn His Pro Lys Ile Glu Thr Ala Ala Met						
	4020		4025			4030
Asp Gly Thr Leu Arg Glu Thr Leu Val Gln Asp Asn Ile Gln Trp Pro						
	4035		4040			4045
Thr Gly Leu Ala Val Asp Tyr His Asn Glu Arg Leu Tyr Trp Ala Asp						
	4050		4055			4060
Ala Lys Leu Ser Val Ile Gly Ser Ile Arg Leu Asn Gly Thr Asp Pro						
	4065		4070			4075
						4080

Ile Val Ala Ala Asp Ser Lys Arg Gly Leu Ser His Pro Phe Ser Ile
 4085 4090 4095
 Asp Val Phe Glu Asp Tyr Ile Tyr Gly Val Thr Tyr Ile Asn Asn Arg
 4100 4105 4110
 Val Phe Lys Ile His Lys Phe Gly His Ser Pro Leu Tyr Asn Leu Thr
 4115 4120 4125
 Gly Gly Leu Ser His Ala Ser Asp Val Val Leu Tyr His Gln His Lys
 4130 4135 4140
 Gln Pro Glu Val Thr Asn Pro Cys Asp Arg Lys Lys Cys Glu Trp Leu
 4145 4150 4155 4160
 Cys Leu Leu Ser Pro Ser Gly Pro Val Cys Thr Cys Pro Asn Gly Lys
 4165 4170 4175
 Arg Leu Asp Asn Gly Thr Cys Val Pro Val Pro Ser Pro Thr Pro Pro
 4180 4185 4190
 Pro Asp Ala Pro Arg Pro Gly Thr Cys Thr Leu Gln Cys Phe Asn Gly
 4195 4200 4205
 Gly Ser Cys Phe Leu Asn Ala Arg Arg Gln Pro Lys Cys Arg Cys Gln
 4210 4215 4220
 Pro Arg Tyr Thr Gly Asp Lys Cys Glu Leu Asp Gln Cys Trp Glu Tyr
 4225 4230 4235 4240
 Cys His Asn Gly Gly Thr Cys Ala Ala Ser Pro Ser Gly Met Pro Thr
 4245 4250 4255
 Cys Arg Cys Pro Thr Gly Phe Thr Gly Pro Lys Cys Thr Ala Gln Val
 4260 4265 4270
 Cys Ala Gly Tyr Cys Ser Asn Asn Ser Thr Cys Thr Val Asn Gln Gly
 4275 4280 4285
 Asn Gln Pro Gln Cys Arg Cys Leu Pro Gly Phe Leu Gly Asp Arg Cys
 4290 4295 4300
 Gln Tyr Arg Gln Cys Ser Gly Phe Cys Glu Asn Phe Gly Thr Cys Gln
 4305 4310 4315 4320
 Met Ala Ala Asp Gly Ser Arg Gln Cys Arg Cys Thr Val Tyr Phe Glu
 4325 4330 4335
 Gly Pro Arg Cys Glu Val Asn Lys Cys Ser Arg Cys Leu Gln Gly Ala
 4340 4345 4350
 Cys Val Val Asn Lys Gln Thr Gly Asp Val Thr Cys Asn Cys Thr Asp
 4355 4360 4365
 Gly Arg Val Ala Pro Ser Cys Leu Thr Cys Ile Asp His Cys Ser Asn
 4370 4375 4380
 Gly Gly Ser Cys Thr Met Asn Ser Lys Met Met Pro Glu Cys Gln Cys
 4385 4390 4395 4400
 Pro Pro His Met Thr Gly Pro Arg Cys Gln Glu Gln Val Val Ser Gln
 4405 4410 4415
 Gln Gln Pro Gly His Met Ala Ser Ile Leu Ile Pro Leu Leu Leu Leu
 4420 4425 4430
 Leu Leu Leu Leu Val Ala Gly Val Val Phe Trp Tyr Lys Arg Arg
 4435 4440 4445
 Val Arg Gly Ala Lys Gly Phe Gln His Gln Arg Met Thr Asn Gly Ala
 4450 4455 4460
 Met Asn Val Glu Ile Gly Asn Pro Thr Tyr Lys Met Tyr Glu Gly Gly
 4465 4470 4475 4480
 Glu Pro Asp Asp Val Gly Gly Leu Leu Asp Ala Asp Phe Ala Leu Asp
 4485 4490 4495
 Pro Asp Lys Pro Thr Asn Phe Thr Asn Pro Val Tyr Ala Thr Leu Tyr
 4500 4505 4510
 Met Gly Gly His Gly Ser Arg His Ser Leu Ala Ser Thr Asp Glu Lys
 4515 4520 4525
 Arg Glu Leu Leu Gly Arg Gly Pro Glu Asp Glu Ile Gly Asp Pro Leu
 4530 4535 4540
 Ala
 4545

<210> 3
 <211> 4577
 <212> DNA
 <213> Homo sapiens

<400> 3

gctacaatcc	atctcgggtctc	ctccagctcc	ttcttttctgc	aacatgggga	agaacaaaact	60
ccttcattcca	agtctgggttc	ttctcctctt	ggtcctcctg	cccacagacg	cctcagttctc	120
tggaaaaccg	cagtatatgg	ttctgggtccc	ctccctgctc	cacactgaga	ccactgagaa	180
gggctgtgtc	cttctgagct	acctgaatga	gacagtgact	gtaagtgtct	ccttggagtc	240
tgtcagggga	aacaggagcc	tcttcactga	cctggaggcg	gagaatgacg	tactccactg	300
tgtcgccctt	gctgtcccaa	agtcttcctc	caatgaggag	gtaatgttcc	tactgttcca	360
agtgaaggga	ccaacccaag	aatttaagaa	gcggaccaca	gtgatgggta	agaacgagga	420
cagtctgggtc	tttgtccaga	cagacaaatc	aatctacaaa	ccagggcaga	cagtgaatt	480
tctgtgtgtc	tccatggatg	aaaactttca	ccccctgaat	gagttgattc	cactagtata	540
cattcaggat	cccaaaggaa	atcgcatcgc	acaatggcag	agtttccagt	tagaggggtg	600
cctcaagcaa	ttttcttttc	ccctctctatc	agagcccttc	cagggtcct	acaagggtgt	660
ggtacagaag	aaatcagggtg	gaaggacaga	gcaccctttc	accgtggagg	aatttgttct	720
tcccaagttt	gaagtacaag	taacagtgcc	aaagataatc	accatcttgg	aagaagagat	780
gaatgtatca	gtgtgtggcc	tatacacata	tgggaagcct	gtccctggac	atgtgactgt	840
gagcattttg	agaaagtata	gtgacgttcc	cgactgccac	gggtgaagatt	cacagggttt	900
ctgtgagaaa	ttcagtggac	agctaaacag	ccatggctgc	ttctatcagc	aagtaaaaaac	960
caagggtcttc	cagctgaaga	ggaaggagta	tgaatgaaa	cttcacactg	aggcccagat	1020
ccaagaagaa	ggaacagtgg	tgggaattgac	tgggaaggcag	tccagtgaag	tcacaagaac	1080
cataaccaaa	ctctcatttg	tgaagtgga	ctcacacttt	cgacagggga	ttcccttctt	1140
tgggcagggtg	cgccctagtag	atgggaaagg	cgctccctata	ccaaataaag	tcatattcat	1200
cagaggaaat	gaagcaaact	attactccaa	tgtaccaccg	gatgagcatg	gccttgtaca	1260
gttctctatc	aacaccacca	acgttatggg	tacctctctt	actgttaggg	tcaattacaa	1320
ggatcgtagt	ccctgttacg	gctaccagtg	ggtgtcagaa	gaacacgaag	aggcatcatca	1380
cactgcttat	cttgtgttct	ccccaagcaa	gagctttgtc	caccttgagc	ccatgtctca	1440
tgaactaccc	tgtggccata	ctcagacagt	ccaggcacat	tatattctga	atggaggcac	1500
cctgctgggg	ctgaagaagc	tctcctttta	ttatctgata	atggcaaagg	gaggcattgt	1560
ccgaactggg	actcatggac	tgttctgtga	gcaggaagac	atgaagggcc	atttttccat	1620
ctcaatccct	gtgaagtcag	acattgctcc	tgtcgtcggg	ttgctcatct	atgctgtttt	1680
acctaccggg	gacgtgattg	gggattctgc	aaaatatgat	gttgaaaatt	gtctggccaa	1740
caagggtggat	ttgagcttca	gcccattaca	aagtctccca	gcctcacacg	cccacctgcg	1800
agtcacagcg	gctcctcagt	ccgtctgcgc	cctccgtgct	gtggaccaaa	gcgtgctgct	1860
cattgaagcct	gatgctgagc	tctcggcgtc	ctcggtttac	aacctgctac	cagaaaagga	1920
cctcactggc	ttccctgggc	ctttgaatga	ccaggacgat	gaagactgca	tcaatcgtca	1980
taatgtctat	attaatggaa	tcacatatac	tccagtatca	agtacaaatg	aaaaggatat	2040
gtacagcttc	ctagaggaca	tgggcttaaa	ggcattcacc	aactcaaaga	ttcgtaaacc	2100
caaaatgtgt	ccacagcttc	aacagtatga	aatgcatgga	cctgaaggct	tacgtgtagg	2160
tttttatgag	tcagatgtaa	tgggaagagg	ccatgcacgc	ctggtgcatg	ttgaagagcc	2220
tcacacggag	accgtacgaa	agtacttccc	tgagacatgg	atctgggatt	tgggtggtgt	2280
aaactcagca	ggggtggctg	aggtaggagt	aacagtcctt	gacaccatca	ccgagtggaa	2340
ggcaggggcc	ttctgcctgt	ctgaagatgc	tggacttggt	atctcttcca	ctgcctctct	2400
ccgagccctt	cagcccttct	ttgtggagct	tacaatgcct	tactctgtga	ttcgtggaga	2460
ggccttcaca	ctcaaggcca	cggctcctaaa	ctaccttccc	aaatgcatcc	gggtcagttg	2520
gcagctggaa	gcctctcccg	ccttctcttg	tgtcccagtg	gagaaggaa	aagcgctca	2580
ctgcatctgt	gcaaacgggc	ggcaaactgt	gtcctgggca	gtaaccccaa	agtcattagg	2640
aaatgtgaat	ttcactgtga	gcgcagaggc	actagagtct	caagagctgt	gtgggactga	2700
ggtgccttca	gttcctgaac	acggaaggaa	agacacagtc	atcaagcctc	tgttggttga	2760
acctgaagga	ctagagaagg	aaacaacatt	caactcccta	ctttgtccat	cagggtggtga	2820
ggtttctgaa	gaattatccc	tgaacttgcc	accaaatgtg	gtagaagaat	ctgcccagac	2880
ttctgtctca	gttttgggag	acatataggc	ctctgccatg	caaaacacac	aaaatcttct	2940
ccagatggcc	tatggctgtg	gagagcagaa	tatggctcct	tttgcctcta	acatctatgt	3000
actggattat	ctaaatgaaa	cacagcagct	tactccagag	gtcaagtcca	aggccattgg	3060
ctatctcaac	actggttacc	agagacagtt	gaactacaaa	cactatgatg	gctcctacag	3120
cacctttggg	gagcgatatg	gcaggaacca	gggcaacacc	tggctcacag	cctttgttct	3180
gaagactttt	gccaagctc	gagcctacat	cttcacgatg	gaagcacaca	ttacccaagc	3240
cctcatatgg	ctctcccaga	ggcagaagga	caatggctgt	ttcaggagct	ctgggtcact	3300

gctcaacaat	gccataaagg	gaggagtaga	agatgaagtg	accctctccg	cctatatcac	3360
catcgccctt	ctggagattc	ctctcacagt	cactcaccct	gttgtccgca	atgccctgtt	3420
ttgacctggag	tcagcctgga	agacagcaca	agaaggggac	catggcagcc	atgtatatac	3480
caaagcactg	ctggcctatg	cttttgcctt	ggcaggtaac	caggacaaga	ggaaggaagt	3540
actcaagtca	cttaatgagg	aagctgtgaa	gaaagacaac	tctgtccatt	gggagcgccc	3600
tcagaaaccc	aaggcaccag	tggggcattt	ttacgaaccc	caggctccct	ctgctgaggt	3660
ggagatgaca	tcctatgtgc	tcctcgctta	tctcacggcc	cagccagccc	caacctcgga	3720
ggacctgacc	tctgcaacca	acatcgtgaa	gtggatcacg	aagcagcaga	atgcccaggg	3780
cggttttctcc	tccacccagg	acacagtggg	ggctctccat	gctctgtcca	aatatggagc	3840
cgccacattt	accaggactg	ggaaggctgc	acaggtgact	atccagtctt	cagggacatt	3900
ttccagcaaa	ttccaagtgg	acaacaacaa	tcgcctgtta	ctgcagcagg	tctcattgcc	3960
agagctgcct	ggggaatata	gcatgaaagt	gacaggagaa	ggatgtgtct	acctccagac	4020
ctccttgaaa	tacaatatct	tcccagaaaa	ggaagagttc	ccctttgctt	taggagtgca	4080
gactctgcct	caaacttgtg	atgaacccaa	agcccacacc	agcttccaaa	tctccctaag	4140
tgtcagttac	acagggagcc	gctctgcctc	caacatggcg	atcgttgatg	tgaagatggg	4200
ctctggcttc	attcccctga	agccaacagt	gaaaatgctt	gaaagatcta	accatgtgag	4260
ccggacagaa	gtcagcagca	accatgtctt	gatttacctt	gataaggtgt	caaatcagac	4320
actgagcttg	ttcttcacgg	ttctgcaaga	tgtcccagta	agagatctca	aaccagccat	4380
agtgaaagtc	tatgattact	acgagacgga	tgagtttgca	atcgctgagt	acaatgctcc	4440
ttgcagcaaa	gatcttgga	atgcttgaag	accacaaggc	tgaaaagtgc	tttgctggag	4500
tctgtttctc	tgagctccac	agaagacacg	tgtttttgta	tctttaaaga	cttgatgaat	4560
aaacactttt	tctggctc					4577

<210> 4
 <211> 4422
 <212> DNA
 <213> Homo sapiens

<400> 4						
atgggggaaga	acaaactcct	tcataccaagt	ctggtttcttc	tcctcttggt	cctcctgccc	60
acagacgcct	cagtctctgg	aaaaccgcag	tatatggttc	tgggtcccctc	cctgctccac	120
actgagacca	ctgagaaggg	ctgtgtcctt	ctgagctacc	tgaatgagac	agtgactgta	180
agtgttctct	tggagtctgt	caggggaaac	aggagcctct	tcactgacct	ggaggcggag	240
aatgacgtac	tccactgtgt	cgcttctcgt	gtcccaaagt	cttcatccaa	tgaggaggta	300
atgttctctca	ctgtccaagt	gaaaggacca	acccaagaat	ttaagaagcg	gaccacagtg	360
atggttaaga	acgaggacag	tctggtcttt	gtccagacag	acaaatcaat	ctacaaacca	420
gggcagacag	tgaaatttcg	tgttgtctcc	atggatgaaa	actttcacc	cctgaatgag	480
ttgattccac	tagtatacat	tcaggatccc	aaaggaaatc	gcatcgca	atggcagagt	540
ttccagttag	agggtggcct	caagcaattt	tcttttcccc	tctcatcaga	gcccttccag	600
ggctcctaca	aggtggtggt	acagaagaaa	tcagggtgga	ggacagagca	ccctttcacc	660
gtggagggaat	ttgttcttcc	caagtttgaa	gtacaagtaa	cagtgcctaa	gataatcacc	720
atcttggaag	aagagatgaa	tgtatcagtg	tgtggcctat	acacatatgg	gaagcctgtc	780
cctggacatg	tgactgtgag	catttgcaga	aagtatagtg	acgcttccga	ctgccacggg	840
gaagattcac	aggctttctg	tgagaaattc	agtggacagc	taaacagcca	tggctgcttc	900
tatcagcaag	taaaaaccaa	ggtcttccag	ctgaagagga	aggagtatga	aatgaaactt	960
cacactgagg	cccagatcca	agaagaagga	acagtgggtg	aattgactgg	aaggcagtcc	1020
agtgaatatca	caagaacat	aaccaaactc	tcatttgtga	aagtggactc	acactttcga	1080
caggggaattc	ccttcttttg	gcagggtgcg	ctagtagatg	ggaaaggcgt	ccctatacca	1140
aataaagtca	tattcatcag	aggaaatgaa	gcaaactatt	actccaatgc	taccacggat	1200
gagcatggcc	ttgtacagtt	ctctatcaac	accaccaacg	ttatgggtac	ctctcttact	1260
gttaggggtca	attacaagga	tcgtagtccc	tgttacggct	accagtgggt	gtcagaagaa	1320
cacgaagagg	cacatcacac	tgcttatctt	gtgttctccc	caagcaagag	ctttgtccac	1380
cttgagccca	tgtctcatga	actaccctgt	ggccatactc	agacagtcca	ggcacattat	1440
attctgaatg	gaggcaccct	gctggggctg	aagaagctct	ccttttatta	tctgataatg	1500
gcaaaggagg	gcattgtccg	aactgggact	catggactgc	ttgtgaagca	ggaagacatg	1560
aagggccatt	tttccatctc	aatccctgtg	aagtcagaca	ttgctcctgt	cgctcggttg	1620
ctcatctatg	ctgtttttacc	taccggggac	gtgattgggg	attctgcaaa	atatgatgtt	1680
gaaaattgtc	tggccaacaa	ggtggatttg	agcttcagcc	catcacaag	tctcccagcc	1740
tcacacgccc	acctgcgagt	cacagcggct	cctcagtcct	tctgcgccct	ccgtgctgtg	1800
gaccaaagcg	tgctgctcat	gaagcctgat	gctgagctct	cggcgtcctc	ggtttacaac	1860
ctgctaccag	aaaaggacct	cactggcttc	cctgggcctt	tgaatgacca	ggacgatgaa	1920

gactgcatca	atcgtcataa	tgtctatatt	aatggaatca	catatactcc	agtatcaagt	1980
acaaatgaaa	aggatatgta	cagcttccta	gaggacatgg	gcttaaaggc	attcaccaac	2040
tcaaagattc	gtaaacccaa	aatgtgtcca	cagcttcaac	agtatgaaat	gcatggacct	2100
gaagggtctac	gtgtaggttt	ttatgagtca	gatgtaatgg	gaagaggcca	tgcacgcctg	2160
gtgcatgttg	aagagcctca	cacggagacc	gtacgaaagt	acttccctga	gacatggatc	2220
tgggatttgg	tgggtgtaaa	ctcagcaggg	gtggctgagg	taggagtaac	agtccctgac	2280
accatcaccg	agtggaaggc	aggggccttc	tgcctgtctg	aagatgctgg	acttgggtatc	2340
tcttccactg	cctctctccg	agccttccag	cccttctttg	tggagcttac	aatgccttac	2400
tctgtgattc	gtggagaggc	cttcacactc	aaggccacgg	tcctaaacta	ccttcccaaa	2460
tgcattccggg	tcagtgtgca	gctggaagcc	tctcccgcct	tccttgctgt	cccagtgagg	2520
aaggaacaag	cgcctcactg	catctgtgca	aacgggcggc	aaactgtgtc	ctgggcagta	2580
accccaaagt	cattaggaaa	tgtgaatttc	actgtgagcg	cagaggcact	agagtctcaa	2640
gagctgtgtg	ggactgaggt	gccttcagtt	cctgaacacg	gaaggaaaga	cacagtcatc	2700
aagcctctgt	tggttgaacc	tgaaggacta	gagaaggaaa	caacattcaa	ctccctactt	2760
tgtccatcag	gtggtgaggt	ttctgaagaa	ttatccctga	aactgccacc	aaatgtggta	2820
gaagaatctg	cccagagcttc	tgtctcagtt	ttgggagaca	tattaggctc	tgccatgcaa	2880
aacacacaaa	atcttctcca	gatgccctat	ggctgtggag	agcagaatat	ggtcctcttt	2940
gctcctaaca	tctatgtact	ggattatcta	aatgaaacac	agcagcttac	tccagaggtc	3000
aagtccaagg	ccattggcta	tctcaacact	ggttaccaga	gacagttgaa	ctacaaacac	3060
tatgatggct	cttcacgacac	ctttggggag	cgatatggca	ggaaccaggg	caacacctgg	3120
ctcagagcct	ttgttctgaa	gacttttgcc	caagctcgag	cctacatctt	catcgatgaa	3180
gcacacatta	cccaagccct	catatggctc	tcccagaggc	agaaggacaa	tggctgtttc	3240
aggagctctg	ggtcactgct	caacaatgcc	ataaagggag	gagtagaaga	tgaagtgacc	3300
ctctccgcct	atatcaccat	cgcccttctg	gagattcctc	tcacagtcac	tcaccctgtt	3360
gtccgcaatg	ccctgttttg	cctggagtc	gcctggaaga	cagcacaaga	aggggaccat	3420
ggcagccatg	tatataccaa	agcactgctg	gcctatgctt	ttgccctggc	aggtaaccag	3480
gacaagagga	aggaagtact	caagtcactt	aatgaggaag	ctgtgaagaa	agacaactct	3540
gtccattggg	agcgccctca	gaaacccaag	gcaccagtgg	ggcattttta	cgaaccccag	3600
gctccctctg	ctgaggtgga	gatgacatcc	tatgtgctcc	tcgcttatct	cacggcccag	3660
ccagccccc	cctcggagga	cctgacctct	gcaaccaaca	tcgtgaagtg	gatcacgaag	3720
cagcagaatg	cccaggggcg	tttctcctcc	acccaggaca	cagtgggtggc	tctccatgct	3780
ctgtccaaat	atggagccgc	cacatttacc	aggactggga	aggctgcaca	ggtgactatc	3840
cagtcttcag	ggacattttc	cagcaaattc	caagtggaca	acaacaatcg	cctgttactg	3900
cagcaggtct	cattgccaga	gctgcctggg	gaatacagca	tgaaagtgac	aggagaagga	3960
tgtgtctacc	tccagacctc	cttgaaatac	aatattctcc	cagaaaagga	agagttcccc	4020
tttgcttttag	gagtgcagac	tctgcctcaa	acttgtgatg	aacccaaagc	ccacaccagc	4080
ttccaaatct	ccctaagtgt	cagttacaca	gggagccgct	ctgcctccaa	catggcgatc	4140
gttgatgtga	agatggcttc	tggcttcatt	ccctggaagc	caacagtga	aatgcttgaa	4200
agatctaacc	atgtgagccg	gacagaagtc	agcagcaacc	atgtcttgat	ttaccttgat	4260
aaggtgtcaa	atcagacact	gagcttgttc	ttcacggttc	tgcaagatgt	cccagtaaga	4320
gatctcaaac	cagccatagt	gaaagtctat	gattactacg	agacggatga	gtttgcaatc	4380
gctgagtaca	atgctccttg	cagcaaagat	cttggaaatg	ct		4422

<210> 5

<211> 1474

<212> PRT

<213> Homo sapiens

<400> 5

Met	Gly	Lys	Asn	Lys	Leu	Leu	His	Pro	Ser	Leu	Val	Leu	Leu	Leu	Leu
1				5					10					15	
Val	Leu	Leu	Pro	Thr	Asp	Ala	Ser	Val	Ser	Gly	Lys	Pro	Gln	Tyr	Met
			20					25					30		
Val	Leu	Val	Pro	Ser	Leu	Leu	His	Thr	Glu	Thr	Thr	Glu	Lys	Gly	Cys
			35				40					45			
Val	Leu	Leu	Ser	Tyr	Leu	Asn	Glu	Thr	Val	Thr	Val	Ser	Ala	Ser	Leu
			50			55					60				
Glu	Ser	Val	Arg	Gly	Asn	Arg	Ser	Leu	Phe	Thr	Asp	Leu	Glu	Ala	Glu
65					70					75				80	
Asn	Asp	Val	Leu	His	Cys	Val	Ala	Phe	Ala	Val	Pro	Lys	Ser	Ser	Ser
			85						90					95	

Asn	Glu	Glu	Val	Met	Phe	Leu	Thr	Val	Gln	Val	Lys	Gly	Pro	Thr	Gln
			100					105					110		
Glu	Phe	Lys	Lys	Arg	Thr	Thr	Val	Met	Val	Lys	Asn	Glu	Asp	Ser	Leu
		115					120					125			
Val	Phe	Val	Gln	Thr	Asp	Lys	Ser	Ile	Tyr	Lys	Pro	Gly	Gln	Thr	Val
		130				135					140				
Lys	Phe	Arg	Val	Val	Ser	Met	Asp	Glu	Asn	Phe	His	Pro	Leu	Asn	Glu
145					150					155					160
Leu	Ile	Pro	Leu	Val	Tyr	Ile	Gln	Asp	Pro	Lys	Gly	Asn	Arg	Ile	Ala
			165						170					175	
Gln	Trp	Gln	Ser	Phe	Gln	Leu	Glu	Gly	Gly	Leu	Lys	Gln	Phe	Ser	Phe
			180					185					190		
Pro	Leu	Ser	Ser	Glu	Pro	Phe	Gln	Gly	Ser	Tyr	Lys	Val	Val	Val	Gln
		195					200					205			
Lys	Lys	Ser	Gly	Gly	Arg	Thr	Glu	His	Pro	Phe	Thr	Val	Glu	Glu	Phe
		210				215					220				
Val	Leu	Pro	Lys	Phe	Glu	Val	Gln	Val	Thr	Val	Pro	Lys	Ile	Ile	Thr
225					230					235					240
Ile	Leu	Glu	Glu	Glu	Met	Asn	Val	Ser	Val	Cys	Gly	Leu	Tyr	Thr	Tyr
				245					250					255	
Gly	Lys	Pro	Val	Pro	Gly	His	Val	Thr	Val	Ser	Ile	Cys	Arg	Lys	Tyr
			260					265					270		
Ser	Asp	Ala	Ser	Asp	Cys	His	Gly	Glu	Asp	Ser	Gln	Ala	Phe	Cys	Glu
		275					280					285			
Lys	Phe	Ser	Gly	Gln	Leu	Asn	Ser	His	Gly	Cys	Phe	Tyr	Gln	Gln	Val
		290				295					300				
Lys	Thr	Lys	Val	Phe	Gln	Leu	Lys	Arg	Lys	Glu	Tyr	Glu	Met	Lys	Leu
305					310					315					320
His	Thr	Glu	Ala	Gln	Ile	Gln	Glu	Glu	Gly	Thr	Val	Val	Glu	Leu	Thr
				325					330					335	
Gly	Arg	Gln	Ser	Ser	Glu	Ile	Thr	Arg	Thr	Ile	Thr	Lys	Leu	Ser	Phe
			340					345					350		
Val	Lys	Val	Asp	Ser	His	Phe	Arg	Gln	Gly	Ile	Pro	Phe	Phe	Gly	Gln
		355					360					365			
Val	Arg	Leu	Val	Asp	Gly	Lys	Gly	Val	Pro	Ile	Pro	Asn	Lys	Val	Ile
		370				375					380				
Phe	Ile	Arg	Gly	Asn	Glu	Ala	Asn	Tyr	Tyr	Ser	Asn	Ala	Thr	Thr	Asp
385					390					395					400
Glu	His	Gly	Leu	Val	Gln	Phe	Ser	Ile	Asn	Thr	Thr	Asn	Val	Met	Gly
				405					410					415	
Thr	Ser	Leu	Thr	Val	Arg	Val	Asn	Tyr	Lys	Asp	Arg	Ser	Pro	Cys	Tyr
			420					425					430		
Gly	Tyr	Gln	Trp	Val	Ser	Glu	Glu	His	Glu	Glu	Ala	His	His	Thr	Ala
		435				440						445			
Tyr	Leu	Val	Phe	Ser	Pro	Ser	Lys	Ser	Phe	Val	His	Leu	Glu	Pro	Met
		450				455					460				
Ser	His	Glu	Leu	Pro	Cys	Gly	His	Thr	Gln	Thr	Val	Gln	Ala	His	Tyr
465					470					475					480
Ile	Leu	Asn	Gly	Gly	Thr	Leu	Leu	Gly	Leu	Lys	Lys	Leu	Ser	Phe	Tyr
				485					490					495	
Tyr	Leu	Ile	Met	Ala	Lys	Gly	Gly	Ile	Val	Arg	Thr	Gly	Thr	His	Gly
			500					505					510		
Leu	Leu	Val	Lys	Gln	Glu	Asp	Met	Lys	Gly	His	Phe	Ser	Ile	Ser	Ile
		515					520					525			
Pro	Val	Lys	Ser	Asp	Ile	Ala	Pro	Val	Ala	Arg	Leu	Leu	Ile	Tyr	Ala
					535						540				
Val	Leu	Pro	Thr	Gly	Asp	Val	Ile	Gly	Asp	Ser	Ala	Lys	Tyr	Asp	Val
545					550					555					560
Glu	Asn	Cys	Leu	Ala	Asn	Lys	Val	Asp	Leu	Ser	Phe	Ser	Pro	Ser	Gln
				565					570					575	
Ser	Leu	Pro	Ala	Ser	His	Ala	His	Leu	Arg	Val	Thr	Ala	Ala	Pro	Gln

Arg Gln Lys Asp Asn Gly Cys Phe Arg Ser Ser Gly Ser Leu Leu Asn
 1075 1080 1085
 Asn Ala Ile Lys Gly Gly Val Glu Asp Glu Val Thr Leu Ser Ala Tyr
 1090 1095 1100
 Ile Thr Ile Ala Leu Leu Glu Ile Pro Leu Thr Val Thr His Pro Val
 1105 1110 1115 1120
 Val Arg Asn Ala Leu Phe Cys Leu Glu Ser Ala Trp Lys Thr Ala Gln
 1125 1130 1135
 Glu Gly Asp His Gly Ser His Val Tyr Thr Lys Ala Leu Leu Ala Tyr
 1140 1145 1150
 Ala Phe Ala Leu Ala Gly Asn Gln Asp Lys Arg Lys Glu Val Leu Lys
 1155 1160 1165
 Ser Leu Asn Glu Glu Ala Val Lys Lys Asp Asn Ser Val His Trp Glu
 1170 1175 1180
 Arg Pro Gln Lys Pro Lys Ala Pro Val Gly His Phe Tyr Glu Pro Gln
 1185 1190 1195 1200
 Ala Pro Ser Ala Glu Val Glu Met Thr Ser Tyr Val Leu Leu Ala Tyr
 1205 1210 1215
 Leu Thr Ala Gln Pro Ala Pro Thr Ser Glu Asp Leu Thr Ser Ala Thr
 1220 1225 1230
 Asn Ile Val Lys Trp Ile Thr Lys Gln Gln Asn Ala Gln Gly Gly Phe
 1235 1240 1245
 Ser Ser Thr Gln Asp Thr Val Val Ala Leu His Ala Leu Ser Lys Tyr
 1250 1255 1260
 Gly Ala Ala Thr Phe Thr Arg Thr Gly Lys Ala Ala Gln Val Thr Ile
 1265 1270 1275 1280
 Gln Ser Ser Gly Thr Phe Ser Ser Lys Phe Gln Val Asp Asn Asn Asn
 1285 1290 1295
 Arg Leu Leu Leu Gln Gln Val Ser Leu Pro Glu Leu Pro Gly Glu Tyr
 1300 1305 1310
 Ser Met Lys Val Thr Gly Glu Gly Cys Val Tyr Leu Gln Thr Ser Leu
 1315 1320 1325
 Lys Tyr Asn Ile Leu Pro Glu Lys Glu Glu Phe Pro Phe Ala Leu Gly
 1330 1335 1340
 Val Gln Thr Leu Pro Gln Thr Cys Asp Glu Pro Lys Ala His Thr Ser
 1345 1350 1355 1360
 Phe Gln Ile Ser Leu Ser Val Ser Tyr Thr Gly Ser Arg Ser Ala Ser
 1365 1370 1375
 Asn Met Ala Ile Val Asp Val Lys Met Val Ser Gly Phe Ile Pro Leu
 1380 1385 1390
 Lys Pro Thr Val Lys Met Leu Glu Arg Ser Asn His Val Ser Arg Thr
 1395 1400 1405
 Glu Val Ser Ser Asn His Val Leu Ile Tyr Leu Asp Lys Val Ser Asn
 1410 1415 1420
 Gln Thr Leu Ser Leu Phe Thr Val Leu Gln Asp Val Pro Val Arg
 1425 1430 1435 1440
 Asp Leu Lys Pro Ala Ile Val Lys Val Tyr Asp Tyr Tyr Glu Thr Asp
 1445 1450 1455
 Glu Phe Ala Ile Ala Glu Tyr Asn Ala Pro Cys Ser Lys Asp Leu Gly
 1460 1465 1470
 Asn Ala

<210> 6
 <211> 14896
 <212> DNA
 <213> Homo sapiens

<400> 6
 cagcgggtgcg agctccaggc ccatgcactg aggaggcgga aacaagggga gccccagag
 ctccatcaag cccctccaa aggtccct acccggtcca cgccccccac cccctccc

60
 120

cgccctccctcc	caattgtgca	tttttgcagc	cggaggcgccg	tccgagatgg	ggctgtgagc	180
ttcgcccggg	gagggggaaa	gagcagcgag	gagtgaagcg	gggggggtggg	gtgaagggtt	240
tggatttcgg	ggcagggggc	gcacccccgt	cagcaggccc	tccccagggg	gctcggaact	300
ctacctcttc	accacggccc	ctgggtgcgt	ttgccgaagg	aaagaataag	aacagagaag	360
gaggaggggg	aaaggaggaa	aagggggacc	ccccaaactgg	gggggggtgaa	ggagagaagt	420
agcaggacca	gaggggaagg	ggctgtctgt	tgcatacagc	cacaccatgc	tgaccccgcc	480
gttgctcctg	ctgctgcccc	tgctctcagc	tctggctcg	gcggctatcg	acgcccctaa	540
gacttgtagc	cccaagcagt	ttgcctgcag	agatcaaata	acctgtatct	caaagggtctg	600
gcggtgtagc	ggtgagagg	actgcccaga	cggatctgac	gaggccccctg	agatttgtcc	660
acagagtaag	gcccagcgat	gccagccaaa	cgagcataac	tgccctgggtg	ctgagctgtg	720
tgttcccatg	tcccgctct	gcaatgggg	ccaggactgc	atggacggct	cagatgagg	780
gccccactgc	cgagagctcc	aaggcaactg	ctctcgctg	ggctgccagc	accatttgtt	840
ccccacactc	gatgggcccc	cctgctactg	caacagcagc	tttcagcttc	aggcagatgg	900
caagacctgc	aaagattttg	atgagtgtct	agtgtacggc	acctgcagcc	agctatgcac	960
caacacagac	ggctccttca	tatgtggctg	tgttgaagga	tacctcctgc	agccggataa	1020
ccgctcctgc	aaggccaaga	acgagccagt	agaccggccc	cctgtgctgt	tgatagccaa	1080
ctcccagaac	atcttggtcca	cgtacctgag	tggggcccag	gtgtctacca	tcacacctac	1140
gagcacgcgg	cagaccacag	ccatggactt	cagctatgcc	aacgagaccg	tatgctgggt	1200
gcatgtttgg	gacagtgtct	ctcagacgca	gctcaagtgt	gcccgcattgc	ctggcctaaa	1260
gggcttcgtg	gatgagcaca	ccatcaacat	ctccctcagt	ctgcaccacg	tggaaacgat	1320
ggccatcgac	tggctgacag	gcaacttcta	ctttgtggat	gacatcgatg	atagatctt	1380
tgtctgcaac	agaaatgggg	acacatgtgt	cacattgcta	gacctggaac	tctacaaccc	1440
caagggcatt	gccctggacc	ctgccatggg	gaaggtgttt	ttcactgact	atgggcagat	1500
cccaaagggtg	gaacgctgtg	acatggatgg	gcagaaccgc	accaagctcg	tcgacagcaa	1560
gattgtgttt	cctcatggca	tcacgctgga	cctggctcagc	cgccctgtct	actgggcaga	1620
tgccatctctg	gactatattg	aagtgggtga	ctatgagggc	aagggcccgcc	agaccatcat	1680
ccagggcattc	ctgattgagc	acctgtacgg	cctgactgtg	tttgagaatt	atctctatgc	1740
caccaactcg	gacaatgcca	atgcccagca	gaagacgagt	gtgatccgtg	tgaaccgctt	1800
taacagcacc	gagtaccagg	ttgtcaccgg	ggtggacaag	ggtgggtgcc	tcacactcta	1860
ccaccagagg	cgtcagcccc	gagtgaggag	ccatgcctgt	gaaaacgacc	agtatgggaa	1920
gccgggtggc	tgctctgaca	tctgcctgct	ggccaacagc	cacaaggcgc	ggacctgccg	1980
ctgccgttcc	ggcttcagcc	tgggcagtga	cgggaagtca	tgcaagaagc	cggagcatga	2040
gctgttcttc	gtgtatggca	agggccggcc	aggcatcatc	cggggcatgg	atatgggggc	2100
caaggtcccg	gatgagcaca	tgatccccat	tgaaaacctc	atgaaccccc	gagccctgga	2160
cttccacgct	gagaccggct	tcactactt	tgccgacacc	accagctacc	tcattggccg	2220
ccagaagatt	gatggcactg	agcgggagac	catcctgaag	gacggcatcc	acaatgtgga	2280
gggtgtggcc	gtggactgga	tgggagacaa	tctgtactgg	acggacgatg	ggcccaaaaa	2340
gacaatcagc	gtggccaggc	tggagaaagc	tgctcagacc	cgcaagactt	taatcgagg	2400
caaatgaca	caccccagg	ctatttgtgt	ggatccactc	aatgggtgga	tgtactggac	2460
agactgggag	gaggacccca	aggacagtgc	gcgtgggcgg	ctggagagg	cgtggatgga	2520
tggtcacac	cgagacatct	ttgtcacctc	caagacagtg	ctttggccca	atgggctaag	2580
cctggacatc	ccggctgggc	gcctctactg	ggtggatgcc	ttctacgacc	gcatacgagc	2640
gatactgctc	aatggcacag	accggaagat	tgtgtatgaa	ggtcctgagc	tgaaccacgc	2700
ctttggcctg	tgtaaccatg	gcaactacct	cttctggact	gagtatcgga	gtggcagtgt	2760
ctaccgcttg	gaacgggggtg	taggagggcg	acccccact	gtgacccttc	tgcgactgta	2820
gcggcccccc	atctttgaga	tccgaatgta	tgatgccag	cagcagcaag	ttggcaccaa	2880
caaatgccg	gtgaacaatg	gcggctgcag	cagcctgtgc	ttggccaccc	ctgggagccg	2940
ccagtgcgcc	tgtgctgagg	accaggtgtt	ggacgcagac	ggcgtcactt	gcttggcgaa	3000
cccatcctac	gtgcctccac	cccagtgcc	gccaggcgag	tttgccctgtg	ccaacagccg	3060
ctgcatccag	gagcgtgga	agtgtgacgg	agacaacgat	tgccctggaca	acagtgatga	3120
ggccccagcc	ctctgccatc	agcacacctg	cccctcggac	cgattcaagt	gcgagaacaa	3180
ccggtgcac	cccaaccgct	ggctctgcga	cggggacaat	gactgtggga	acagtgaaga	3240
tgagtccaat	gccacttgtt	cagcccgcac	ctgccccccc	aaccagttct	cctgtgccag	3300
tggccgctgc	atccccatct	cctggacgtg	tgatctggat	gacgactgtg	gggaccgctc	3360
tgatgagtct	gcttcgtgtg	cctatcccac	ctgcttcccc	ctgactcagt	ttacctgcaa	3420
caatggcaga	tgtatcaaca	tcaactggag	atgcgacaat	gacaatgact	gtggggacaa	3480
cagtgcagaa	gccggctgca	gccactcctg	ttctagcacc	cagttcaagt	gcaacagcgg	3540
gcgttgcatc	cccagcact	ggacctgcga	tggggacaat	gactgcggag	actacagtga	3600
tgagacacac	gccaactgca	ccaaccaggc	cacgaggccc	cctggtggct	gccacactga	3660
tgagttccag	tgccggctgg	atggactatg	catccccctg	cggtggcgct	gcgatgggga	3720
cactgactgc	atggactcca	gcgatgagaa	gagctgtgag	ggagtgaacc	acgtctgcga	3780

tcccagtgtc	aagtttggct	gcaaggactc	agctcggtgc	atcagcaaag	cgtgggtgtg	3840
tgatggcgac	aatgactgtg	aggataactc	ggacgaggag	aactgcgagt	ccctggcctg	3900
caggccaccc	tcgcaccctt	gtgccaaaca	cacctcagtc	tgcttgcccc	ctgacaagct	3960
gtgtgatggc	aacgacgact	gtggcgacgg	ctcagatgag	ggcgagctct	gcgaccagtg	4020
ctctctgaat	aacggtggct	gcagccacaa	ctgctcagtg	gcacctggcg	aaggcattgt	4080
gtgttcctgc	cctctgggca	tggagctggg	gcccgcacaac	cacacctgcc	agatccagag	4140
ctactgtgcc	aagcatctca	aatgcagcca	aaagtgcgac	cagaacaagt	tcagcgtgaa	4200
gtgctcctgc	tacgagggct	gggtcctgga	acctgacggc	gagagctgcc	gcagcctgga	4260
ccccttcaag	ccgttcatca	ttttctccaa	ccgccatgaa	atccggcgca	tcgatcttca	4320
caaaggagac	tacagcgtec	tggtgcccgg	cctgcgcaac	accatcgccc	tggacttcca	4380
cctcagccag	agcgccctct	actggaccga	cgtggtggag	gacaagatct	accgcgggaa	4440
gctgctggag	aacggagccc	tgactagtct	cgaggtgggtg	attcagtatg	gcctggccac	4500
acccgagggc	ctggctgtag	actggattgc	aggcaacatc	tactgggtgg	agagtaacct	4560
ggatcagatc	gaggtggcca	agctggatgg	gacctccgg	accacctgc	tggccggtga	4620
cattgagcac	ccaagggcaa	tcgactgga	tccccgggat	gggatcctgt	tttgacaga	4680
ctgggatgcc	agcctgcccc	gcattgaggc	agcctccatg	agtggggctg	ggcgccgcac	4740
cgtgcaccgg	gagaccggct	ctgggggctg	gcccacggg	ctcaccgtgg	actacctgga	4800
gaagcgcac	ctttggattg	acgccaggtc	agatgccatt	tactcagccc	gttacgacgg	4860
ctctggccac	atggaggtgc	ttcggggaca	cgagttcctg	tcgcaccctg	ttgcagtgc	4920
gctgtacggg	ggggaggtct	actggactga	ctggcgaaac	aacacactgg	ctaaggccaa	4980
caagtggacc	ggccacaatg	tcaccgtggt	acagaggacc	aacacccagc	cctttgacct	5040
gcaggtgtac	cacccctccc	gccagcccat	ggctcccaat	ccctgtgagg	ccaatggggg	5100
ccagggcccc	tgctccacc	tgtgtctcat	caactacaac	cggaccgtgt	cctgcgcctg	5160
ccccacctc	atgaagctcc	acaaggacaa	caccacctgc	tatgagtta	agaagttcct	5220
gctgtacgca	cgtcagatgg	agatccgagg	tgtggacctg	gatgctccct	actacaacta	5280
catcatctcc	ttcacggtgc	ccgacatcga	caacgtcaca	gtgctagact	acgatgcccg	5340
cgagcagcgt	gtgtactgg	ctgacgtgcg	gacacaggcc	atcaagcggg	ccttcatcaa	5400
cggcacaggc	gtggagacag	tcgtctctgc	agacttgcca	aatgcccacg	ggctggctgt	5460
ggatgagggt	tcccgaaacc	tgttctggac	aagctatgac	accaataaga	agcagatcaa	5520
tgtggcccgg	ctggatggct	ccttcaagaa	cgcagtgggtg	cagggcctgg	agcagcccca	5580
tggccttgtc	gtccaccctc	tgcgtgggaa	gctctactgg	accgatgggtg	acaacatcag	5640
catggccaac	atggatggca	gcaatcgcac	cctgctcttc	agtggccaga	agggccccgt	5700
gggcctggct	attgacttcc	ctgaaagcaa	actctactgg	atcagctccg	ggaaccatac	5760
catcaaccgc	tgcaacctgg	atgggagtg	gctggaggtc	atcgatgcca	tgcgagacca	5820
gctgggcaag	gccaccgccc	tggccatcat	gggggacaag	ctgtggtggg	ctgatcaggt	5880
gtcggaaaag	atgggcacat	gcagcaaggc	tgacggctcg	ggctccgtgg	tccttcggaa	5940
cagcaccacc	ctggtgatgc	acatgaaggt	ctatgacgag	agcatccagc	tggaccataa	6000
gggcaccaac	ccctgcagtg	tcaacaacgg	tgactgtctc	cagctctgcc	tgcccacgtc	6060
agagacgacc	cgctcctgca	tgtgcacagc	cggctatagc	ctccggagtg	gccagcaggc	6120
ctgcgagggc	gtaggttcc	ttctcctgta	ctctgtgcat	gagggaatca	ggggaattcc	6180
cctggatccc	aatgacaagt	cagatgccct	ggctcccagt	tccgggacct	cgctggctgt	6240
cggcatcgac	ttccacgctg	aaaatgacac	catctactgg	gtggacatgg	gcctgagcac	6300
gatcagccgg	gccaagcggg	accagacgtg	gcgtgaagac	gtggtgacca	atggcattgg	6360
ccgtgtggag	ggcattgcag	tggactggat	cgcaggcaac	atctactgga	cagaccaggg	6420
ctttgatgtc	atcgaggtcg	ccgggtcaa	tggctccttc	cgctacgtgg	tgatctccca	6480
gggtctagac	aagccccggg	ccatcacctg	ccaccggag	aaagggtact	tgttctggac	6540
tgagtggggt	cagtatccgc	gtattgagcg	gtctcggcta	gatggcacgg	agcgtgtggt	6600
gctgggtcaac	gtcagcatca	gctggcccaa	cggcatctca	gtggactacc	aggatgggaa	6660
gctgtactgg	tgcatgcac	ggacagacaa	gattgaacgg	atcgacctgg	agacaggtga	6720
gaaccgcgag	gtggttctgt	ccagcaacaa	catggacatg	ttttcagtgt	ctgtgtttga	6780
ggatttcatc	tactggagtg	acaggactca	tgccaacggc	tctatcaagc	gcgggagcaa	6840
agacaatgcc	acagactccg	tgcccctgcg	aaccggcatc	ggcgtccagc	ttaaagacat	6900
caaagtcttc	aaccgggacc	ggcagaaaag	caccaacgtg	tgcgcggtgg	ccaatggcgg	6960
gtgccagcag	ctgtgacctg	accggggccg	tgggcagcgg	gcctgcgcct	gtgcccacgg	7020
gatgctggct	gaagacggag	catcgtgcgc	cgagtatgcc	ggctacctgc	tctactcaga	7080
gcgcaccatt	ctcaagagta	tccacctgcg	ggatgagcgc	aacctcaatg	cgcccgtgca	7140
gcccttcgag	gacctgagc	acatgaagaa	cgtcatcgcc	ctggcctttg	actaccgggc	7200
aggcacctct	ccgggcaccc	ccaatcgcat	cttcttcagc	gacatccact	ttgggaacat	7260
ccaacagatc	aacgacgatg	gctccaggag	gatcaccatt	gtggaaaacg	tgggctccgt	7320
ggaaggcctg	gcctatcacc	gtggctggga	cactctctat	tggacaagct	acacgacatc	7380
caccatcacg	cgccacacag	tggaccagac	cgcgccaggg	gccttcgagc	gtgagaccgt	7440

catcactatg	tetggagatg	accacccacg	ggccttcggt	ttggacgagt	gccagaacct	7500
catgttctgg	accaactgga	atgagcagca	tcccagcadc	atgcgggcg	cgctctcggg	7560
agccaatgtc	ctgaccctta	tcgagaagga	catccgtacc	cccaatggcc	tggccatcga	7620
ccaccgtgcc	gagaagctct	acttctctga	cgccaccctg	gacaagatcg	agcgggtgcga	7680
gtatgacggc	tcccaccgct	atgtgatcct	aaagtccagag	cctgtccacc	ccttcgggct	7740
ggcgtgtat	ggggagcaca	ttttctggac	tgactgggtg	cgccgggcag	tgcagcgggc	7800
caacaagcac	gtgggcagca	acatgaagct	gctgcgcgtg	gacatcccc	agcagcccat	7860
gggcatcatc	gccgtggcca	acgacaccaa	cagctgtgaa	ctctctccat	gccgaatcaa	7920
caacggtggc	tgccaggacc	tgtgtctgct	cactcaccag	ggccatgtca	actgctcatg	7980
cggagggggc	cgaatcctcc	aggatgacct	cacctgccga	gcgggtgaatt	cctcttgccg	8040
agcacaagat	gagtttgagt	gtgccaatgg	cgagtgcadc	aacttcagcc	tgacctgcga	8100
cggcgctccc	cactgcaagg	acaagtccga	tgagaagcca	tcctactgca	actcccgccg	8160
ctgcaagaag	actttccggc	agtgcagcaa	tgggcgctgt	gtgtccaaca	tgtgtgggtg	8220
caacggggcc	gacgactgtg	gggatggctc	tgacgagatc	ccttgcaaca	agacagcctg	8280
tgggtgtggc	gagttccgct	gccgggacgg	gacctgcadc	gggaactcca	gccgctgcaa	8340
ccagtttgtg	gattgtgagg	acgcctcaga	tgagatgaac	tgcagtgcc	ccgactgcag	8400
cagctacttc	cgccctggcg	tgaaggcggt	gctcttccag	ccctgcgagc	ggacctcact	8460
ctgctacgca	cccagctggg	tgtgtgatgg	cgccaatgac	tgtggggact	acagtgatga	8520
gcgcgactgc	ccaggtgtga	aacgccccag	atgcctctcg	aattacttcg	cctgccctag	8580
tgggcgctgc	atccccatga	gctggacgtg	tgacaaagag	gatgactgtg	aacatggcga	8640
ggagcagacc	cactgcaaca	agtctgtctc	agaggcccag	tttgagtgcc	agaacctcgc	8700
ctgcatctcc	aagcagtggc	tgtgtgacgg	cagcgatgac	tgtggggatg	gctcagacga	8760
ggctgctcac	tgtgaaggca	agacgtgcgg	cccctcctcc	ttctcctgcc	ctggcaccga	8820
cgtgtgcgtc	cccagagcgt	ggctctgtga	cggtgacaaa	gactgtgctg	atgggtgcaga	8880
cgagagcatc	gcagctgggt	gcttgtacaa	cagcacttgt	gacgaccgtg	agttcatgtg	8940
ccagaaccgc	cagtgcaccc	ccaagcactt	cgtgtgtgac	cacgaccgtg	actgtgcaga	9000
tggctctgat	gagtcctccg	agtgtgagta	cccagacctg	ggccccagtg	agttccgctg	9060
tgccaatggg	cgctgtctga	gctcccgcga	gtgggagtg	gatggcgaga	atgactgcca	9120
cgaccaagat	gacgaggctc	ccaagaaccc	acactgcacc	agcccagagc	acaagtgcga	9180
tgcctcgta	cagttcctgt	gcagcagtg	gcgctgtgtg	gctgaggcac	tgtctgcaa	9240
cggccaggat	gactgtggcg	acagctcgga	cgagcgtggc	tgccacatca	atgagtgtct	9300
cagccgcaag	ctcagtggct	gcagccagga	ctgtgaggac	ctcaagatcg	gcttcaagtg	9360
ccgctgtgc	cctggcttcc	ggctgaagga	tgacggccgg	acgtgtgctg	atgtggacga	9420
gtgcagcacc	accttcccct	gcagccagcg	ctgcatcaac	acccatggca	gctataagtg	9480
tctgtgtgtg	gagggctatg	caccccgcg	cggcgacccc	cacagctgca	aggctgtgac	9540
tgacgaggaa	ccgtttctga	tcttcgcca	ccggtactac	ctgcgcaagc	tcaacctgga	9600
cgggtccaac	tacacgttac	ttaagcagg	cctgaacaac	gccgttgctt	tggattttga	9660
ctaccgagag	cagatgatct	actggacaga	tgtgaccacc	cagggcagca	tgtatccgaag	9720
gatgcacctt	aacgggagca	atgtgcaggt	cctacacogt	acaggcctca	gcaacccgca	9780
tgggctggct	gtggactggg	tgggtggcaa	cctgtactgg	tgcgacaaag	gccgggacac	9840
catcgagggtg	tccaagctca	atggggccta	tggacgggtg	ctggctcagct	ctggcctccg	9900
tgagcccagg	gctctgtgtg	tggatgtgca	gaatgggtac	ctgtactgga	cagactgggg	9960
tgaccattca	ctgatcggcc	gcacccggcat	ggatgggtcc	agccgcagcg	tcatcgtgga	10020
caccaagatc	acatggccca	atggcctgac	gctggactat	gtcactgagc	gcactactctg	10080
ggccgacgcc	cgcgaggact	acattgaatt	tgccagcctg	gatggctcca	atcgccacgt	10140
tgtgtctgagc	caggacatcc	cgcacatctt	tgactgacc	ctgtttgagg	actacgtcta	10200
ctggaccgac	tgggaaacaa	agtccattaa	ccgagcccac	aagaccacgg	gcaccaacaa	10260
aacgtctctc	atcagcacgc	tgcaccggcc	catggacctg	catgtcttcc	atgcctctgcg	10320
ccagccagac	gtgcccaatc	acccctgcaa	ggtcaacaat	ggtggctgca	gcaacctgtg	10380
cctgtgtgtc	cccgggggag	ggcacaatg	tgcctgcccc	accaacttct	acctgggcag	10440
cgatggggcg	acctgtgtgt	ccaactgcac	ggctagccag	tttgtatgca	agaacgacaa	10500
gtgcatcccc	ttctggtgga	agtgtgacac	cgaggacgac	tgcggggacc	actcagacga	10560
gcccccgga	tgcctgagt	tcaagtgcgg	gccccgacag	ttccagtgtc	ccacaggtat	10620
ctgcacaaa	cctgccttca	tctgcgatgg	cgacaatgac	tgccaggaca	acagtgcaga	10680
ggccaactgt	gacatccacg	tctgcttgcc	cagtcagttc	aaatgcacca	acaccaaccg	10740
ctgtattccc	ggcatcttcc	gctgcaatgg	cgaggacaac	tgcggagatg	gggaggatga	10800
gagggactgc	cccagaggtga	cctgcgcccc	caaccagttc	cagtgtctcca	ttaccaaacg	10860
gtgcatcccc	cgggtctggg	tctgcgaccg	ggacaatgac	tgtgtggatg	gcagtgatga	10920
gccccccaac	tgcaccacga	tgacctgtgg	tgtggacgag	ttccgctgca	aggattcggg	10980
ccgctgcadc	ccagcgcgtt	ggaagtgtga	cggagaggat	gactgtgggg	atggctcgga	11040
tgagcccaag	gaagagtgtg	atgaacgcac	ctgtgagcca	taccagttcc	gctgcaagaa	11100

caaccgctgc	gtgcccggcc	gctggcagtg	cgactacgac	aacgattgcg	gtgacaactc	11160
cgatgaagag	agctggcacc	ctcggccctg	ctccgagagt	gagttctcct	gtgccaacgg	11220
ccgctgcate	gcggggcgct	ggaaatgcga	tggagaccac	gactgcgcgg	acggctcgga	11280
cgagaaagac	tgcaccccc	gctgtgacat	ggaccagttc	cagtgcgaaga	gcggccactg	11340
catccccctg	cgctggcgct	gtgacgcaga	cgccgactgc	atggacggca	gcgacgagga	11400
ggcctgcggc	actggcgctg	ggacctgccc	cctggacgag	ttccagtgc	acaacacctt	11460
gtgcaagccg	ctggcctgga	agtgcgatgg	cgaggatgac	tgtggggaca	actcagatga	11520
gaaccccgag	gagtgtgccc	ggttcgtgtg	ccctcccaac	cggcccttcc	gttgcaagaa	11580
tgaccgcgtc	tgtctgtgga	tcggggcgcca	atgcgatggc	acggacaact	gtggggatgg	11640
gactgatgaa	gaggactgtg	agccccccac	agccccacac	acccactgca	aagacaagaa	11700
ggagtttctg	tgccggaacc	agcgtctgct	ctcctcctcc	ctgcgctgca	acatgttcga	11760
tgactgcggg	gacggctctg	acgaggagga	ctgcagcatc	gaccccaagc	tgaccagctg	11820
cgccaccaat	gccagcatct	gtggggacga	ggcacgctgc	gtgcgcaccg	agaaagcggc	11880
ctactgtgcc	tgccgctcgg	gcttccacac	cgtgcccggc	cagcccggat	gccaagacat	11940
caacgagtg	ctgcgcttcg	gcacctgtct	ccagctctgc	aacaacacca	agggcggcca	12000
cctctgcagc	tgcgctcgga	acttcatgaa	gacgcacaac	acctgcaagg	ccgaaggctc	12060
tgagtaccag	gtcctgtaca	tcgctgatga	caatgagatc	cgcagcctgt	ttcccggcca	12120
ccccattcgg	gcttacgagc	aggcattcca	gggtgacgag	agtgtccgca	ttgatgctat	12180
ggatgtccat	gtcaaggctg	gccgtgtcta	tggaccaac	tggcacacgg	gcaccatctc	12240
ctaccgcgag	ctgccacctc	ctgcgcctcc	taccgattcc	aaccgccacc	ggcgacagat	12300
tgaccggggg	gtcaccacc	tcaacatttc	agggctgaag	atgccagag	gcatcgccat	12360
cgactgggtg	gccggaaacg	tgtactggac	cgactcgggc	cgagatgtga	ttgaggtggc	12420
gcagatgaag	ggcgagaacc	gcaagacgct	catctcgggc	atgattgacg	agccccacgc	12480
cattgtggtg	gacccactga	gggggacat	gtactgggtc	gactggggca	accaccccaa	12540
gattgagacg	gcagcgatgg	atgggacgct	tcgggagaca	ctggtgcagg	acaacattca	12600
gtggcccaca	ggcctggccg	tggattatca	caatgagcgg	ctgtactggg	cagacgcca	12660
gctttcagtc	atcggcagca	tcgggctcaa	tggcacggac	cccatttgtg	ctgctgacag	12720
caaacgaggg	ctaagtcacc	ccttcagcat	cgacgtcttt	gaggattaca	tctatggtgt	12780
cacctacatg	aataatcgct	tcttcaagat	ccataagttt	ggccacagcc	ccttgggtcaa	12840
cctgacaggg	ggcctgagcc	acgcctctga	cgtggctcct	taccatcagc	acaagcagcc	12900
cgaagtgaac	aacccatgtg	accgcaagaa	atgcgagtg	ctctgcctgc	tgagccccag	12960
tgggcctgtc	tgcacctgtc	ccaatgggaa	gcggctggac	aacggcacat	gcgtgcctgt	13020
gcccctctca	acgccccccc	cagatgtctc	ccggcctgga	acctgtaacc	tgacgtgctt	13080
caacggtggc	agctgtttcc	tcaatgcacg	gaggcagccc	aagtgccgct	gccaacccc	13140
ctacacgggt	gacaagtgtg	aactggacca	gtgctgggag	cactgtcgca	atgggggac	13200
ctgtgctgcc	ttcccctctg	gcctgcccac	gtgcgggtgc	cccacgggct	tcacggggcc	13260
caaatgcacc	cagcaggtgt	gtgcgggcta	ctgtgccaac	aacagcacct	gcactgtcaa	13320
ccagggcaac	cagccccagt	gccgatgcct	accggcttcc	ctgggcgacc	gctgccagta	13380
ccggcagtg	tctggctact	gtgagaactt	tggcacatgc	cagatggctg	ctgatggctc	13440
ccgacaatgc	cgctgcactg	cctactttga	gggatcgagg	tgtgaggtga	acaagtgcag	13500
ccgctgtctc	gaaggggcct	gtgtggtcaa	caagcagagt	ggggatgtca	cctgcaactg	13560
cacggatggc	cgggtggccc	ccagctgtct	gacctgcgtc	ggccactgca	gcaatggcgg	13620
ctcctgtacc	atgaacagca	aaatgatgcc	tgagtgccag	tgccccaccc	acatgacagg	13680
gccccgggtg	gaggagcacg	tcttcagcca	gcagcagcca	ggacatatag	cctccatcct	13740
aatccctctg	ctgttgctgc	tgctgctggt	tctggtggcc	ggagtggat	tctggtataa	13800
gcggcgagtc	caaggggcta	agggcttcca	gcaccaacgg	atgaccaacg	gggccatgaa	13860
cgtggagatt	ggaaacccca	cctacaagat	gtacgaaggc	ggagagcctg	atgatgtggg	13920
aggcctactg	gacgctgact	ttgccctgga	ccctgacaag	cccaccaact	tcaccaaccc	13980
cgtgtatgcc	acactctaca	tggggggcca	tggcagtcgc	cactccctgg	ccagcacgga	14040
cgagaagcga	gaactcctgg	gccggggccc	tgaggacgag	ataggggacc	ccttggcata	14100
gggcccctgc	ccgtcggact	gccccagaa	agcctcctgc	cccctgccgg	tgaagtcctt	14160
cagtgaagccc	ctccccagcc	agcccttccc	tggccccgcc	ggatgtataa	atgtaaaaat	14220
gaaggaatta	cattttatat	gtgagcgagc	aagccggcaa	gcgagcacag	tattatttct	14280
ccatcccctc	cctgcctgct	ccttggcacc	cccatgtctc	cttcaggagg	acaggcaggg	14340
agggcttggg	gctgcacctc	ctacctccc	accagaacgc	accccaactg	gagagctggg	14400
gggtcagcct	tcccctccct	gtataagaca	ccttgcgaag	gctctcccct	ctcgccccat	14460
ccctgcttgc	ccgctcccac	agcttctctga	gggctaattc	tgggaaggga	gagttctttg	14520
ctgcccctgt	ctggaagacg	tggctctggg	tgaggtaggc	gggaaaggat	ggagtgtttt	14580
agttcttggg	ggaggccacc	ccaaacccca	gccccaaact	caggggcacc	tatgagatgg	14640
ccatgctcaa	ccccctccc	agacaggccc	tcctgtctc	cagggccccc	accgaggttc	14700
ccagggtctg	agacttcttc	tggtaaacad	tcctccagcc	tcctctcccc	tggggacgcc	14760


```

aaggaggtgg gccacacca ggaagggaaa gcgggcagcc ccgttttggg gacgtgaacg 14820
ttttaataat ttttgctgaa ttctttacaa ctaaataaca cagatattct tataaataaa 14880
attgtaaaaa aaaaaa 14896

```

```

<210> 7
<211> 126
<212> PRT
<213> Homo sapiens

```

```

<400> 7
Ile Ala Leu Asp Phe His Leu Ser Gln Ser Ala Leu Tyr Trp Thr Asp
 1          5          10          15
Val Val Glu Asp Lys Ile Tyr Arg Gly Lys Leu Leu Asp Asn Gly Ala
 20          25          30
Leu Thr Ser Phe Glu Val Val Ile Gln Tyr Gly Leu Ala Thr Pro Glu
 35          40          45
Gly Leu Ala Val Asp Trp Ile Ala Gly Asn Ile Tyr Trp Val Glu Ser
 50          55          60
Asn Leu Asp Gln Ile Glu Val Ala Lys Leu Asp Gly Thr Leu Arg Thr
 65          70          75          80
Thr Leu Leu Ala Gly Asp Ile Glu His Pro Arg Ala Ile Ala Leu Asp
 85          90          95
Pro Arg Asp Gly Ile Leu Phe Trp Thr Asp Trp Asp Ala Ser Leu Pro
100          105          110
Arg Ile Glu Ala Ala Ser Met Ser Gly Ala Gly Arg Arg Thr
115          120          125

```

```

<210> 8
<211> 153
<212> PRT
<213> Homo sapiens

```

```

<400> 8
Leu Leu Gln Gln Val Ser Leu Pro Glu Leu Pro Gly Glu Tyr Ser Met
 1          5          10          15
Lys Val Thr Gly Glu Gly Cys Val Tyr Leu Gln Thr Ser Leu Lys Tyr
 20          25          30
Asn Ile Leu Pro Glu Lys Glu Glu Phe Pro Phe Ala Leu Gly Val Gln
 35          40          45
Thr Leu Pro Gln Thr Cys Asp Glu Pro Lys Ala His Thr Ser Phe Gln
 50          55          60
Ile Ser Leu Ser Val Ser Tyr Thr Gly Ser Arg Ser Ala Ser Asn Met
 65          70          75          80
Ala Ile Val Asp Val Lys Met Val Ser Gly Phe Ile Pro Leu Lys Pro
 85          90          95
Thr Val Lys Met Leu Glu Arg Ser Asn His Val Ser Arg Thr Glu Val
100          105          110
Ser Ser Asn His Val Leu Ile Tyr Leu Asp Lys Val Ser Asn Gln Thr
115          120          125
Leu Ser Leu Phe Phe Thr Val Leu Gln Asp Val Pro Val Arg Asp Leu
130          135          140
Lys Pro Ala Ile Val Lys Val Tyr Asp
145          150

```

```

<210> 9
<211> 138
<212> PRT
<213> Homo sapiens

```

<400> 9
 Met Lys Val Thr Gly Glu Gly Cys Val Tyr Leu Gln Thr Ser Leu Lys
 1 5 10 15
 Tyr Asn Ile Leu Pro Glu Lys Glu Glu Phe Pro Phe Ala Leu Gly Val
 20 25 30
 Gln Thr Leu Pro Gln Thr Cys Asp Glu Pro Lys Ala His Thr Ser Phe
 35 40 45
 Gln Ile Ser Leu Ser Val Ser Tyr Thr Gly Ser Arg Ser Ala Ser Asn
 50 55 60
 Met Ala Ile Val Asp Val Lys Met Val Ser Gly Phe Ile Pro Leu Lys
 65 70 75 80
 Pro Thr Val Lys Met Leu Glu Arg Ser Asn His Val Ser Arg Thr Glu
 85 90 95
 Val Ser Ser Asn His Val Leu Ile Tyr Leu Asp Lys Val Ser Asn Gln
 100 105 110
 Thr Leu Ser Leu Phe Phe Thr Val Leu Gln Asp Val Pro Val Arg Asp
 115 120 125
 Leu Lys Pro Ala Ile Val Lys Val Tyr Asp
 130 135

<210> 10
 <211> 27
 <212> PRT
 <213> Homo sapiens

<400> 10
 Ser Val Ser Tyr Thr Gly Ser Arg Ser Ala Ser Asn Met Ala Ile Val
 1 5 10 15
 Asp Val Lys Met Val Ser Gly Phe Ile Pro Leu
 20 25

<210> 11
 <211> 126
 <212> PRT
 <213> Homo sapiens

<400> 11
 Leu Gln Gln Val Ser Leu Pro Glu Leu Pro Gly Glu Tyr Ser Met Lys
 1 5 10 15
 Val Thr Gly Glu Gly Cys Val Tyr Leu Gln Thr Ser Leu Lys Tyr Asn
 20 25 30
 Ile Leu Pro Glu Lys Glu Glu Phe Pro Phe Ala Leu Gly Val Gln Thr
 35 40 45
 Leu Pro Gln Thr Cys Asp Glu Pro Lys Ala His Thr Ser Phe Gln Ile
 50 55 60
 Ser Leu Ser Val Ser Tyr Thr Gly Ser Arg Ser Ala Ser Asn Met Ala
 65 70 75 80
 Ile Val Asp Val Lys Met Val Ser Gly Phe Ile Pro Leu Lys Pro Thr
 85 90 95
 Val Lys Met Leu Glu Arg Ser Asn His Val Ser Arg Thr Glu Val Ser
 100 105 110
 Ser Asn His Val Leu Ile Tyr Leu Asp Lys Val Ser Asn Gln
 115 120 125

<210> 12
 <211> 111
 <212> PRT
 <213> Homo sapiens

<400> 12
 Leu Gln Gln Val Ser Leu Pro Glu Leu Pro Gly Glu Tyr Ser Met Lys

1		5		10		15									
Val	Thr	Gly	Glu	Gly	Cys	Val	Tyr	Leu	Gln	Thr	Ser	Leu	Lys	Tyr	Asn
		20						25					30		
Ile	Leu	Pro	Glu	Lys	Glu	Glu	Phe	Pro	Phe	Ala	Leu	Gly	Val	Gln	Thr
		35					40					45			
Leu	Pro	Gln	Thr	Cys	Asp	Glu	Pro	Lys	Ala	His	Thr	Ser	Phe	Gln	Ile
		50				55					60				
Ser	Leu	Ser	Val	Ser	Tyr	Thr	Gly	Ser	Arg	Ser	Ala	Ser	Asn	Met	Ala
65					70					75					80
Ile	Val	Asp	Val	Lys	Met	Val	Ser	Gly	Phe	Ile	Pro	Leu	Lys	Pro	Thr
				85					90					95	
Val	Lys	Met	Leu	Glu	Arg	Ser	Asn	His	Val	Ser	Arg	Thr	Glu	Val	
			100					105					110		

<210> 13
 <211> 81
 <212> PRT
 <213> Homo sapiens

	<400> 13														
Leu	Gln	Gln	Val	Ser	Leu	Pro	Glu	Leu	Pro	Gly	Glu	Tyr	Ser	Met	Lys
1				5					10					15	
Val	Thr	Gly	Glu	Gly	Cys	Val	Tyr	Leu	Gln	Thr	Ser	Leu	Lys	Tyr	Asn
		20						25					30		
Ile	Leu	Pro	Glu	Lys	Glu	Glu	Phe	Pro	Phe	Ala	Leu	Gly	Val	Gln	Thr
		35					40					45			
Leu	Pro	Gln	Thr	Cys	Asp	Glu	Pro	Lys	Ala	His	Thr	Ser	Phe	Gln	Ile
		50				55					60				
Ser	Leu	Ser	Val	Ser	Tyr	Thr	Gly	Ser	Arg	Ser	Ala	Ser	Asn	Met	Ala
65					70					75					80
Ile															

<210> 14
 <211> 101
 <212> PRT
 <213> Homo sapiens

	<400> 14														
Gln	Thr	Ser	Leu	Lys	Tyr	Asn	Ile	Leu	Pro	Glu	Lys	Glu	Glu	Phe	Pro
1				5					10					15	
Phe	Ala	Leu	Gly	Val	Gln	Thr	Leu	Pro	Gln	Thr	Cys	Asp	Glu	Pro	Lys
		20						25					30		
Ala	His	Thr	Ser	Phe	Gln	Ile	Ser	Leu	Ser	Val	Ser	Tyr	Thr	Gly	Ser
		35					40					45			
Arg	Ser	Ala	Ser	Asn	Met	Ala	Ile	Val	Asp	Val	Lys	Met	Val	Ser	Gly
		50				55					60				
Phe	Ile	Pro	Leu	Lys	Pro	Thr	Val	Lys	Met	Leu	Glu	Arg	Ser	Asn	His
65					70					75					80
Val	Ser	Arg	Thr	Glu	Val	Ser	Ser	Asn	His	Val	Leu	Ile	Tyr	Leu	Asp
				85					90					95	
Lys	Val	Ser	Asn	Gln											
			100												

<210> 15
 <211> 76
 <212> PRT
 <213> Homo sapiens

	<400> 15														
Gln	Thr	Ser	Leu	Lys	Tyr	Asn	Ile	Leu	Pro	Glu	Lys	Glu	Glu	Phe	Pro

1		5		10		15									
Phe	Ala	Leu	Gly	Val	Gln	Thr	Leu	Pro	Gln	Thr	Cys	Asp	Glu	Pro	Lys
			20					25					30		
Ala	His	Thr	Ser	Phe	Gln	Ile	Ser	Leu	Ser	Val	Ser	Tyr	Thr	Gly	Ser
		35					40					45			
Arg	Ser	Ala	Ser	Asn	Met	Ala	Ile	Val	Asp	Val	Lys	Met	Val	Ser	Gly
	50					55					60				
Phe	Ile	Pro	Leu	Lys	Pro	Thr	Val	Lys	Met	Leu	Glu				
65					70					75					

<210> 16
 <211> 56
 <212> PRT
 <213> Homo sapiens

	<400> 16														
Gln	Thr	Ser	Leu	Lys	Tyr	Asn	Ile	Leu	Pro	Glu	Lys	Glu	Glu	Phe	Pro
1			5					10						15	
Phe	Ala	Leu	Gly	Val	Gln	Thr	Leu	Pro	Gln	Thr	Cys	Asp	Glu	Pro	Lys
		20						25					30		
Ala	His	Thr	Ser	Phe	Gln	Ile	Ser	Leu	Ser	Val	Ser	Tyr	Thr	Gly	Ser
		35					40					45			
Arg	Ser	Ala	Ser	Asn	Met	Ala	Ile								
	50					55									

<210> 17
 <211> 76
 <212> PRT
 <213> Homo sapiens

	<400> 17														
Gln	Thr	Cys	Asp	Glu	Pro	Lys	Ala	His	Thr	Ser	Phe	Gln	Ile	Ser	Leu
1			5					10						15	
Ser	Val	Ser	Tyr	Thr	Gly	Ser	Arg	Ser	Ala	Ser	Asn	Met	Ala	Ile	Val
		20					25						30		
Asp	Val	Lys	Met	Val	Ser	Gly	Phe	Ile	Pro	Leu	Lys	Pro	Thr	Val	Lys
		35					40					45			
Met	Leu	Glu	Arg	Ser	Asn	His	Val	Ser	Arg	Thr	Glu	Val	Ser	Ser	Asn
	50					55					60				
His	Val	Leu	Ile	Tyr	Leu	Asp	Lys	Val	Ser	Asn	Gln				
65					70					75					

<210> 18
 <211> 76
 <212> PRT
 <213> Homo sapiens

	<400> 18														
Gln	Thr	Cys	Asp	Glu	Pro	Lys	Ala	His	Thr	Ser	Phe	Gln	Ile	Ser	Leu
1			5					10						15	
Ser	Val	Ser	Tyr	Thr	Gly	Ser	Arg	Ser	Ala	Ser	Asn	Met	Ala	Ile	Val
		20					25						30		
Asp	Val	Lys	Met	Val	Ser	Gly	Phe	Ile	Pro	Leu	Lys	Pro	Thr	Val	Lys
		35					40					45			
Met	Leu	Glu	Arg	Ser	Asn	His	Val	Ser	Arg	Thr	Glu	Val	Ser	Ser	Asn
	50					55					60				
His	Val	Leu	Ile	Tyr	Leu	Asp	Lys	Val	Ser	Asn	Gln				
65					70					75					

<210> 19
 <211> 31

<212> PRT
 <213> Homo sapiens

<400> 19
 Gln Thr Cys Asp Glu Pro Lys Ala His Thr Ser Phe Gln Ile Ser Leu
 1 5 10 15
 Ser Val Ser Tyr Thr Gly Ser Arg Ser Ala Ser Asn Met Ala Ile
 20 25 30

<210> 20
 <211> 44
 <212> PRT
 <213> Homo sapiens

<400> 20
 Lys Thr Cys Ser Pro Lys Gln Phe Ala Cys Arg Asp Gln Ile Thr Cys
 1 5 10 15
 Ile Ser Lys Gly Trp Arg Cys Asp Gly Glu Arg Asp Cys Pro Asp Gly
 20 25 30
 Ser Asp Glu Ala Pro Glu Ile Cys Pro Gln Ser Lys
 35 40

<210> 21
 <211> 86
 <212> PRT
 <213> Homo sapiens

<400> 21
 Lys Thr Cys Ser Pro Lys Gln Phe Ala Cys Arg Asp Gln Ile Thr Cys
 1 5 10 15
 Ile Ser Lys Gly Trp Arg Cys Asp Gly Glu Arg Asp Cys Pro Asp Gly
 20 25 30
 Ser Asp Glu Ala Pro Glu Ile Cys Pro Gln Ser Lys Ala Gln Arg Cys
 35 40 45
 Gln Pro Asn Glu His Asn Cys Leu Gly Thr Glu Leu Cys Val Pro Met
 50 55 60
 Ser Arg Leu Cys Asn Gly Val Gln Asp Cys Met Asp Gly Ser Asp Glu
 65 70 75 80
 Gly Pro His Cys Arg Glu
 85

<210> 22
 <211> 43
 <212> PRT
 <213> Homo sapiens

<400> 22
 Lys Ala Gln Arg Cys Gln Pro Asn Glu His Asn Cys Leu Gly Thr Glu
 1 5 10 15
 Leu Cys Val Pro Met Ser Arg Leu Cys Asn Gly Val Gln Asp Cys Met
 20 25 30
 Asp Gly Ser Asp Glu Gly Pro His Cys Arg Glu
 35 40

<210> 23
 <211> 42
 <212> PRT
 <213> Homo sapiens

<400> 23
 Gln Cys Gln Pro Gly Glu Phe Ala Cys Ala Asn Ser Arg Cys Ile Gln

1		5		10		15									
Glu	Arg	Trp	Lys	Cys	Asp	Gly	Asp	Asn	Asp	Cys	Leu	Asp	Asn	Ser	Asp
			20					25					30		
Glu	Ala	Pro	Ala	Leu	Cys	His	Gln	His	Thr						
		35					40								

<210> 24
 <211> 82
 <212> PRT
 <213> Homo sapiens

<400> 24															
Gln	Cys	Gln	Pro	Gly	Glu	Phe	Ala	Cys	Ala	Asn	Ser	Arg	Cys	Ile	Gln
1				5					10					15	
Glu	Arg	Trp	Lys	Cys	Asp	Gly	Asp	Asn	Asp	Cys	Leu	Asp	Asn	Ser	Asp
			20					25					30		
Glu	Ala	Pro	Ala	Leu	Cys	His	Gln	His	Thr	Cys	Pro	Ser	Asp	Arg	Phe
		35					40					45			
Lys	Cys	Glu	Asn	Asn	Arg	Cys	Ile	Pro	Asn	Arg	Trp	Leu	Cys	Asp	Gly
	50					55					60				
Asp	Asn	Asp	Cys	Gly	Asn	Ser	Glu	Asp	Glu	Ser	Asn	Ala	Thr	Cys	Ser
65					70					75					80
Ala	Arg														

<210> 25
 <211> 122
 <212> PRT
 <213> Homo sapiens

<400> 25															
Gln	Cys	Gln	Pro	Gly	Glu	Phe	Ala	Cys	Ala	Asn	Ser	Arg	Cys	Ile	Gln
1				5					10					15	
Glu	Arg	Trp	Lys	Cys	Asp	Gly	Asp	Asn	Asp	Cys	Leu	Asp	Asn	Ser	Asp
			20					25					30		
Glu	Ala	Pro	Ala	Leu	Cys	His	Gln	His	Thr	Cys	Pro	Ser	Asp	Arg	Phe
		35					40					45			
Lys	Cys	Glu	Asn	Asn	Arg	Cys	Ile	Pro	Asn	Arg	Trp	Leu	Cys	Asp	Gly
	50					55					60				
Asp	Asn	Asp	Cys	Gly	Asn	Ser	Glu	Asp	Glu	Ser	Asn	Ala	Thr	Cys	Ser
65					70					75					80
Ala	Arg	Thr	Cys	Pro	Pro	Asn	Gln	Phe	Ser	Cys	Ala	Ser	Gly	Arg	Cys
				85					90					95	
Ile	Pro	Ile	Ser	Trp	Thr	Cys	Asp	Leu	Asp	Asp	Asp	Cys	Gly	Asp	Arg
			100					105					110		
Ser	Asp	Glu	Ser	Ala	Ser	Cys	Ala	Tyr	Pro						
		115					120								

<210> 26
 <211> 161
 <212> PRT
 <213> Homo sapiens

<400> 26															
Gln	Cys	Gln	Pro	Gly	Glu	Phe	Ala	Cys	Ala	Asn	Ser	Arg	Cys	Ile	Gln
1				5					10					15	
Glu	Arg	Trp	Lys	Cys	Asp	Gly	Asp	Asn	Asp	Cys	Leu	Asp	Asn	Ser	Asp
			20					25					30		
Glu	Ala	Pro	Ala	Leu	Cys	His	Gln	His	Thr	Cys	Pro	Ser	Asp	Arg	Phe
		35					40					45			
Lys	Cys	Glu	Asn	Asn	Arg	Cys	Ile	Pro	Asn	Arg	Trp	Leu	Cys	Asp	Gly

50	55	60
Asp Asn Asp Cys Gly	Asn Ser Glu Asp Glu Ser	Asn Ala Thr Cys Ser
65	70	75
Ala Arg Thr Cys Pro	Pro Asn Gln Phe Ser Cys Ala Ser Gly Arg Cys	80
	85	90
Ile Pro Ile Ser Trp Thr Cys Asp	Leu Asp Asp Asp Cys Gly Asp Arg	95
	100	105
Ser Asp Glu Ser Ala Ser Cys Ala Tyr Pro Thr Cys Phe	Pro Leu Thr	110
	115	120
Gln Phe Thr Cys Asn Asn Gly Arg Cys Ile Asn Ile Asn Trp Arg Cys		125
	130	135
Asp Asn Asp Asn Asp Cys Gly Asp Asn Ser Asp Glu Ala Gly Cys Ser		140
145	150	155
His		160

<210> 27
 <211> 208
 <212> PRT
 <213> Homo sapiens

<400> 27
Gln Cys Gln Pro Gly Glu Phe Ala Cys Ala Asn Ser Arg Cys Ile Gln
1 5 10 15
Glu Arg Trp Lys Cys Asp Gly Asp Asn Asp Cys Leu Asp Asn Ser Asp
20 25 30
Glu Ala Pro Ala Leu Cys His Gln His Thr Cys Pro Ser Asp Arg Phe
35 40 45
Lys Cys Glu Asn Asn Arg Cys Ile Pro Asn Arg Trp Leu Cys Asp Gly
50 55 60
Asp Asn Asp Cys Gly Asn Ser Glu Asp Glu Ser Asn Ala Thr Cys Ser
65 70 75 80
Ala Arg Thr Cys Pro Pro Asn Gln Phe Ser Cys Ala Ser Gly Arg Cys
85 90 95
Ile Pro Ile Ser Trp Thr Cys Asp Leu Asp Asp Asp Cys Gly Asp Arg
100 105 110
Ser Asp Glu Ser Ala Ser Cys Ala Tyr Pro Thr Cys Phe Pro Leu Thr
115 120 125
Gln Phe Thr Cys Asn Asn Gly Arg Cys Ile Asn Ile Asn Trp Arg Cys
130 135 140
Asp Asn Asp Asn Asp Cys Gly Asp Asn Ser Asp Glu Ala Gly Cys Ser
145 150 155 160
His Ser Cys Ser Ser Thr Gln Phe Lys Cys Asn Ser Gly Arg Cys Ile
165 170 175
Pro Glu His Trp Thr Cys Asp Gly Asp Asn Asp Cys Gly Asp Tyr Ser
180 185 190
Asp Glu Thr His Ala Asn Cys Thr Asn Gln Ala Thr Arg Pro Pro Gly
195 200 205

<210> 28
 <211> 150
 <212> PRT
 <213> Homo sapiens

<400> 28
Gln Cys Gln Pro Gly Glu Phe Ala Cys Ala Asn Ser Arg Cys Ile Gln
1 5 10 15
Glu Arg Trp Lys Cys Asp Gly Asp Asn Asp Cys Leu Asp Asn Ser Asp
20 25 30
Glu Ala Pro Ala Leu Cys His Gln His Thr Cys Pro Ser Asp Arg Phe
35 40 45

Lys	Cys	Glu	Asn	Asn	Arg	Cys	Ile	Pro	Asn	Arg	Trp	Leu	Cys	Asp	Gly
50						55					60				
Asp	Asn	Asp	Cys	Gly	Asn	Ser	Glu	Asp	Glu	Ser	Asn	Ala	Thr	Cys	Ser
65					70					75					80
Ala	Arg	Thr	Cys	Pro	Pro	Asn	Gln	Phe	Ser	Cys	Ala	Ser	Gly	Arg	Cys
				85					90					95	
Ile	Pro	Ile	Ser	Trp	Thr	Cys	Asp	Leu	Asp	Asp	Asp	Cys	Gly	Asp	Arg
			100					105					110		
Ser	Asp	Glu	Ser	Ala	Ser	Cys	Ala	Tyr	Pro	Thr	Cys	Phe	Pro	Leu	Thr
		115					120					125			
Gln	Phe	Thr	Cys	Asn	Asn	Gly	Arg	Cys	Ile	Asn	Ile	Asn	Trp	Arg	Cys
	130					135					140				
Asp	Asn	Asp	Asn	Asp	Cys										
145					150										

<210> 29
 <211> 231
 <212> PRT
 <213> Homo sapiens

Gln	Cys	Gln	Pro	Gly	Glu	Phe	Ala	Cys	Ala	Asn	Ser	Arg	Cys	Ile	Gln
1				5				10						15	
Glu	Arg	Trp	Lys	Cys	Asp	Gly	Asp	Asn	Asp	Cys	Leu	Asp	Asn	Ser	Asp
			20					25					30		
Glu	Ala	Pro	Ala	Leu	Cys	His	Gln	His	Thr	Cys	Pro	Ser	Asp	Arg	Phe
		35				40						45			
Lys	Cys	Glu	Asn	Asn	Arg	Cys	Ile	Pro	Asn	Arg	Trp	Leu	Cys	Asp	Gly
	50					55					60				
Asp	Asn	Asp	Cys	Gly	Asn	Ser	Glu	Asp	Glu	Ser	Asn	Ala	Thr	Cys	Ser
65					70					75					80
Ala	Arg	Thr	Cys	Pro	Pro	Asn	Gln	Phe	Ser	Cys	Ala	Ser	Gly	Arg	Cys
				85					90					95	
Ile	Pro	Ile	Ser	Trp	Thr	Cys	Asp	Leu	Asp	Asp	Asp	Cys	Gly	Asp	Arg
			100					105					110		
Ser	Asp	Glu	Ser	Ala	Ser	Cys	Ala	Tyr	Pro	Thr	Cys	Phe	Pro	Leu	Thr
		115					120					125			
Gln	Phe	Thr	Cys	Asn	Asn	Gly	Arg	Cys	Ile	Asn	Ile	Asn	Trp	Arg	Cys
	130					135					140				
Asp	Asn	Asp	Asn	Asp	Cys	Gly	Asp	Asn	Ser	Asp	Glu	Ala	Gly	Cys	Ser
145					150					155				160	
His	Ser	Cys	Ser	Ser	Thr	Gln	Phe	Lys	Cys	Asn	Ser	Gly	Arg	Cys	Ile
			165					170						175	
Pro	Glu	His	Trp	Thr	Cys	Asp	Gly	Asp	Asn	Asp	Cys	Gly	Asp	Tyr	Ser
			180					185					190		
Asp	Glu	Thr	His	Ala	Asn	Cys	Thr	Asn	Gln	Ala	Thr	Arg	Pro	Pro	Gly
		195					200					205			
Gly	Cys	His	Thr	Asp	Glu	Phe	Gln	Cys	Arg	Leu	Asp	Gly	Leu	Cys	Ile
	210					215					220				
Pro	Leu	Arg	Trp	Arg	Cys	Asp									
225					230										

<210> 30
 <211> 40
 <212> PRT
 <213> Homo sapiens

Cys	Pro	Ser	Asp	Arg	Phe	Lys	Cys	Glu	Asn	Asn	Arg	Cys	Ile	Pro	Asn
1				5				10						15	
Arg	Trp	Leu	Cys	Asp	Gly	Asp	Asn	Asp	Cys	Gly	Asn	Ser	Glu	Asp	Glu

20 25 30
 Ser Asn Ala Thr Cys Ser Ala Arg
 35 40
 <210> 31
 <211> 80
 <212> PRT
 <213> Homo sapiens
 <400> 31
 Cys Pro Ser Asp Arg Phe Lys Cys Glu Asn Asn Arg Cys Ile Pro Asn
 1 5 10 15
 Arg Trp Leu Cys Asp Gly Asp Asn Asp Cys Gly Asn Ser Glu Asp Glu
 20 25 30
 Ser Asn Ala Thr Cys Ser Ala Arg Thr Cys Pro Pro Asn Gln Phe Ser
 35 40 45
 Cys Ala Ser Gly Arg Cys Ile Pro Ile Ser Trp Thr Cys Asp Leu Asp
 50 55 60
 Asp Asp Cys Gly Asp Arg Ser Asp Glu Ser Ala Ser Cys Ala Tyr Pro
 65 70 75 80

<210> 32
 <211> 119
 <212> PRT
 <213> Homo sapiens

<400> 32
 Cys Pro Ser Asp Arg Phe Lys Cys Glu Asn Asn Arg Cys Ile Pro Asn
 1 5 10 15
 Arg Trp Leu Cys Asp Gly Asp Asn Asp Cys Gly Asn Ser Glu Asp Glu
 20 25 30
 Ser Asn Ala Thr Cys Ser Ala Arg Thr Cys Pro Pro Asn Gln Phe Ser
 35 40 45
 Cys Ala Ser Gly Arg Cys Ile Pro Ile Ser Trp Thr Cys Asp Leu Asp
 50 55 60
 Asp Asp Cys Gly Asp Arg Ser Asp Glu Ser Ala Ser Cys Ala Tyr Pro
 65 70 75 80
 Thr Cys Phe Pro Leu Thr Gln Phe Thr Cys Asn Asn Gly Arg Cys Ile
 85 90 95
 Asn Ile Asn Trp Arg Cys Asp Asn Asp Asn Asp Cys Gly Asp Asn Ser
 100 105 110
 Asp Glu Ala Gly Cys Ser His
 115

<210> 33
 <211> 166
 <212> PRT
 <213> Homo sapiens

<400> 33
 Cys Pro Ser Asp Arg Phe Lys Cys Glu Asn Asn Arg Cys Ile Pro Asn
 1 5 10 15
 Arg Trp Leu Cys Asp Gly Asp Asn Asp Cys Gly Asn Ser Glu Asp Glu
 20 25 30
 Ser Asn Ala Thr Cys Ser Ala Arg Thr Cys Pro Pro Asn Gln Phe Ser
 35 40 45
 Cys Ala Ser Gly Arg Cys Ile Pro Ile Ser Trp Thr Cys Asp Leu Asp
 50 55 60
 Asp Asp Cys Gly Asp Arg Ser Asp Glu Ser Ala Ser Cys Ala Tyr Pro
 65 70 75 80
 Thr Cys Phe Pro Leu Thr Gln Phe Thr Cys Asn Asn Gly Arg Cys Ile

				85					90					95			
Asn	Ile	Asn	Trp	Arg	Cys	Asp	Asn	Asp	Asn	Asp	Cys	Gly	Asp	Asn	Ser		
			100					105					110				
Asp	Glu	Ala	Gly	Cys	Ser	His	Ser	Cys	Ser	Ser	Thr	Gln	Phe	Lys	Cys		
		115					120					125					
Asn	Ser	Gly	Arg	Cys	Ile	Pro	Glu	His	Trp	Thr	Cys	Asp	Gly	Asp	Asn		
	130					135					140						
Asp	Cys	Gly	Asp	Tyr	Ser	Asp	Glu	Thr	His	Ala	Asn	Cys	Thr	Asn	Gln		
145					150					155					160		
Ala	Thr	Arg	Pro	Pro	Gly												
				165													

<210> 34
 <211> 108
 <212> PRT
 <213> Homo sapiens

Cys	Pro	Ser	Asp	Arg	Phe	Lys	Cys	Glu	Asn	Asn	Arg	Cys	Ile	Pro	Asn		
1				5					10					15			
Arg	Trp	Leu	Cys	Asp	Gly	Asp	Asn	Asp	Cys	Gly	Asn	Ser	Glu	Asp	Glu		
		20						25					30				
Ser	Asn	Ala	Thr	Cys	Ser	Ala	Arg	Thr	Cys	Pro	Pro	Asn	Gln	Phe	Ser		
		35				40						45					
Cys	Ala	Ser	Gly	Arg	Cys	Ile	Pro	Ile	Ser	Trp	Thr	Cys	Asp	Leu	Asp		
	50					55					60						
Asp	Asp	Cys	Gly	Asp	Arg	Ser	Asp	Glu	Ser	Ala	Ser	Cys	Ala	Tyr	Pro		
65					70					75					80		
Thr	Cys	Phe	Pro	Leu	Thr	Gln	Phe	Thr	Cys	Asn	Asn	Gly	Arg	Cys	Ile		
				85					90					95			
Asn	Ile	Asn	Trp	Arg	Cys	Asp	Asn	Asp	Asn	Asp	Cys						
			100					105									

<210> 35
 <211> 289
 <212> PRT
 <213> Homo sapiens

Cys	Pro	Ser	Asp	Arg	Phe	Lys	Cys	Glu	Asn	Asn	Arg	Cys	Ile	Pro	Asn		
1				5					10					15			
Arg	Trp	Leu	Cys	Asp	Gly	Asp	Asn	Asp	Cys	Gly	Asn	Ser	Glu	Asp	Glu		
		20						25					30				
Ser	Asn	Ala	Thr	Cys	Ser	Ala	Arg	Thr	Cys	Pro	Pro	Asn	Gln	Phe	Ser		
		35				40						45					
Cys	Ala	Ser	Gly	Arg	Cys	Ile	Pro	Ile	Ser	Trp	Thr	Cys	Asp	Leu	Asp		
	50					55					60						
Asp	Asp	Cys	Gly	Asp	Arg	Ser	Asp	Glu	Ser	Ala	Ser	Cys	Ala	Tyr	Pro		
65					70					75					80		
Thr	Cys	Phe	Pro	Leu	Thr	Gln	Phe	Thr	Cys	Asn	Asn	Gly	Arg	Cys	Ile		
				85					90					95			
Asn	Ile	Asn	Trp	Arg	Cys	Asp	Asn	Asp	Asn	Asp	Cys	Gly	Asp	Asn	Ser		
			100					105					110				
Asp	Glu	Ala	Gly	Cys	Ser	His	Ser	Cys	Ser	Ser	Thr	Gln	Phe	Lys	Cys		
		115					120					125					
Asn	Ser	Gly	Arg	Cys	Ile	Pro	Glu	His	Trp	Thr	Cys	Asp	Gly	Asp	Asn		
	130					135					140						
Asp	Cys	Gly	Asp	Tyr	Ser	Asp	Glu	Thr	His	Ala	Asn	Cys	Thr	Asn	Gln		
145					150					155					160		
Ala	Thr	Arg	Pro	Pro	Gly	Gly	Cys	His	Thr	Asp	Glu	Phe	Gln	Cys	Arg		
				165					170					175			

Leu Asp Gly Leu Cys Ile Pro Leu Arg Trp Arg Cys Asp Gly Asp Thr
 180 185 190
 Asp Cys Met Asp Ser Ser Asp Glu Lys Ser Cys Glu Gly Val Thr His
 195 200 205
 Val Cys Asp Pro Ser Val Lys Phe Gly Cys Lys Asp Ser Ala Arg Cys
 210 215 220
 Ile Ser Lys Ala Trp Val Cys Asp Gly Asp Asn Asp Cys Glu Asp Asn
 225 230 235 240
 Ser Asp Glu Glu Asn Cys Glu Ser Leu Ala Cys Arg Pro Pro Ser His
 245 250 255
 Pro Cys Ala Asn Asn Thr Ser Val Cys Leu Pro Pro Asp Lys Leu Cys
 260 265 270
 Asp Gly Asn Asp Asp Cys Gly Asp Gly Ser Asp Glu Gly Glu Leu Cys
 275 280 285
 Asp

<210> 36
 <211> 40
 <212> PRT
 <213> Homo sapiens

<400> 36
 Thr Cys Pro Pro Asn Gln Phe Ser Cys Ala Ser Gly Arg Cys Ile Pro
 1 5 10 15
 Ile Ser Trp Thr Cys Asp Leu Asp Asp Asp Cys Gly Asp Arg Ser Asp
 20 25 30
 Glu Ser Ala Ser Cys Ala Tyr Pro
 35 40

<210> 37
 <211> 79
 <212> PRT
 <213> Homo sapiens

<400> 37
 Thr Cys Pro Pro Asn Gln Phe Ser Cys Ala Ser Gly Arg Cys Ile Pro
 1 5 10 15
 Ile Ser Trp Thr Cys Asp Leu Asp Asp Asp Cys Gly Asp Arg Ser Asp
 20 25 30
 Glu Ser Ala Ser Cys Ala Tyr Pro Thr Cys Phe Pro Leu Thr Gln Phe
 35 40 45
 Thr Cys Asn Asn Gly Arg Cys Ile Asn Ile Asn Trp Arg Cys Asp Asn
 50 55 60
 Asp Asn Asp Cys Gly Asp Asn Ser Asp Glu Ala Gly Cys Ser His
 65 70 75

<210> 38
 <211> 126
 <212> PRT
 <213> Homo sapiens

<400> 38
 Thr Cys Pro Pro Asn Gln Phe Ser Cys Ala Ser Gly Arg Cys Ile Pro
 1 5 10 15
 Ile Ser Trp Thr Cys Asp Leu Asp Asp Asp Cys Gly Asp Arg Ser Asp
 20 25 30
 Glu Ser Ala Ser Cys Ala Tyr Pro Thr Cys Phe Pro Leu Thr Gln Phe
 35 40 45
 Thr Cys Asn Asn Gly Arg Cys Ile Asn Ile Asn Trp Arg Cys Asp Asn
 50 55 60

Asp	Asn	Asp	Cys	Gly	Asp	Asn	Ser	Asp	Glu	Ala	Gly	Cys	Ser	His	Ser
65				70					75					80	
Cys	Ser	Ser	Thr	Gln	Phe	Lys	Cys	Asn	Ser	Gly	Arg	Cys	Ile	Pro	Glu
				85				90					95		
His	Trp	Thr	Cys	Asp	Gly	Asp	Asn	Asp	Cys	Gly	Asp	Tyr	Ser	Asp	Glu
			100				105						110		
Thr	His	Ala	Asn	Cys	Thr	Asn	Gln	Ala	Thr	Arg	Pro	Pro	Gly		
		115					120					125			

<210> 39
 <211> 68
 <212> PRT
 <213> Homo sapiens

Thr	Cys	Pro	Pro	Asn	Gln	Phe	Ser	Cys	Ala	Ser	Gly	Arg	Cys	Ile	Pro
1				5				10						15	
Ile	Ser	Trp	Thr	Cys	Asp	Leu	Asp	Asp	Asp	Cys	Gly	Asp	Arg	Ser	Asp
			20				25						30		
Glu	Ser	Ala	Ser	Cys	Ala	Tyr	Pro	Thr	Cys	Phe	Pro	Leu	Thr	Gln	Phe
		35				40						45			
Thr	Cys	Asn	Asn	Gly	Arg	Cys	Ile	Asn	Ile	Asn	Trp	Arg	Cys	Asp	Asn
	50					55					60				
Asp	Asn	Asp	Cys												
65															

<210> 40
 <211> 248
 <212> PRT
 <213> Homo sapiens

Cys	Pro	Pro	Asn	Gln	Phe	Ser	Cys	Ala	Ser	Gly	Arg	Cys	Ile	Pro	Ile
1			5					10					15		
Ser	Trp	Thr	Cys	Asp	Leu	Asp	Asp	Asp	Cys	Gly	Asp	Arg	Ser	Asp	Glu
			20				25						30		
Ser	Ala	Ser	Cys	Ala	Tyr	Pro	Thr	Cys	Phe	Pro	Leu	Thr	Gln	Phe	Thr
		35				40					45				
Cys	Asn	Asn	Gly	Arg	Cys	Ile	Asn	Ile	Asn	Trp	Arg	Cys	Asp	Asn	Asp
	50				55					60					
Asn	Asp	Cys	Gly	Asp	Asn	Ser	Asp	Glu	Ala	Gly	Cys	Ser	His	Ser	Cys
65				70					75					80	
Ser	Ser	Thr	Gln	Phe	Lys	Cys	Asn	Ser	Gly	Arg	Cys	Ile	Pro	Glu	His
			85					90					95		
Trp	Thr	Cys	Asp	Gly	Asp	Asn	Asp	Cys	Gly	Asp	Tyr	Ser	Asp	Glu	Thr
			100				105						110		
His	Ala	Asn	Cys	Thr	Asn	Gln	Ala	Thr	Arg	Pro	Pro	Gly	Gly	Cys	His
		115				120						125			
Thr	Asp	Glu	Phe	Gln	Cys	Arg	Leu	Asp	Gly	Leu	Cys	Ile	Pro	Leu	Arg
	130				135						140				
Trp	Arg	Cys	Asp	Gly	Asp	Thr	Asp	Cys	Met	Asp	Ser	Ser	Asp	Glu	Lys
145				150					155					160	
Ser	Cys	Glu	Gly	Val	Thr	His	Val	Cys	Asp	Pro	Ser	Val	Lys	Phe	Gly
			165					170					175		
Cys	Lys	Asp	Ser	Ala	Arg	Cys	Ile	Ser	Lys	Ala	Trp	Val	Cys	Asp	Gly
		180					185					190			
Asp	Asn	Asp	Cys	Glu	Asp	Asn	Ser	Asp	Glu	Glu	Asn	Cys	Glu	Ser	Leu
	195					200						205			
Ala	Cys	Arg	Pro	Pro	Ser	His	Pro	Cys	Ala	Asn	Asn	Thr	Ser	Val	Cys
	210				215						220				
Leu	Pro	Pro	Asp	Lys	Leu	Cys	Asp	Gly	Asn	Asp	Asp	Cys	Gly	Asp	Gly

Asp Gly Asn Asp Asp Cys Gly Asp Gly Ser Asp Glu Gly Glu Leu Cys
65 70 75 80
Asp

<210> 53
<211> 40
<212> PRT
<213> Homo sapiens

<400> 53
Ala Cys Arg Pro Ser His Pro Cys Ala Asn Asn Thr Ser Val Cys
1 5 10 15
Leu Pro Pro Asp Lys Leu Cys Asp Gly Asn Asp Asp Cys Gly Asp Gly
20 25 30
Ser Asp Glu Gly Glu Leu Cys Asp
35 40

<210> 54
<211> 10
<212> PRT
<213> Homo sapiens

<400> 54
Ser Gly Phe Ser Leu Gly Ser Asp Gly Lys
1 5 10

<210> 55
<211> 10
<212> PRT
<213> Homo sapiens

<400> 55
Gly Ile Ala Leu Asp Pro Ala Met Gly Lys
1 5 10

<210> 56
<211> 10
<212> PRT
<213> Homo sapiens

<400> 56
Gly Gly Ala Leu His Ile Tyr His Gln Arg
1 5 10

<210> 57
<211> 11
<212> PRT
<213> Homo sapiens

<400> 57
Val Phe Phe Thr Asp Tyr Gly Gln Ile Pro Lys
1 5 10

<210> 58
<211> 9
<212> PRT
<213> Homo sapiens

<400> 58
 Gly Ala Leu His Ile Tyr His Gln Arg
 1 5

<210> 59
 <211> 19
 <212> PRT
 <213> Homo sapiens

<400> 59
 Arg Val Thr Tyr His Ser Pro Ser Tyr Val Tyr His Gln Phe Glu Arg
 1 5 10 15
 Arg Ala Lys